



SEQUENCE LISTING

B20

<110> Watkins, Maren
Olivera, Baldomero M.
Hillyard, David R.
McIntosh, J. Michael
Jones, Robert M.

<120> Alpha-Conotoxin Peptides

<130> 2314-179.A

<140> US 09/493,795
<141> 2000-01-28

<150> US 60/118,381
<151> 1999-01-29

<160> 404

<170> PatentIn Ver. 2.0

<210> 1
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<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Alpha-Conotoxin
Peptide Generic Formula I

<220>
<221> SITE
<222> (1)..(3)
<223> Xaa at residue 1 is des-Xaa, Ile, Leu or Val; Xaa
at residue 2 is des-Xaa, Ala or Gly; Xaa at
residue 3 is des-Xaa, Gly, Trp (D or L), neo-Trp,
halo-Trp or any unnatural aromatic amino acid.

<220>
<221> SITE
<222> (4)..(5)
<223> N-methyl-Lys, Xaa at residue 4 is des-Xaa, Gly,
Trp (D or L), neo-Trp, halo-Trp or any unnatural
aromatic amino acid; Xaa at residue 5 is Glu,
gamma-carboxy-Glu (Gla), Asp, Ala, Thr, Ser, Gly,

<220>
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<222> (5)..(8)
<223> Ile, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any
unnatural hydroxy containing amino acid; Xaa at
residue 8 is Ser, Thr, Arg, ornithine,

<220>
<221> SITE
<222> (8)..(9)
<223> homoarginine, Lys, N,N-dimethyl-Lys, N,N,N-
trimethyl-Lys or any unnatural basic amino acid;
Xaa at residue 9 is Asp, Glu, Gla, Arg, ornithine,
homoarginine, Lys, N-methyl-Lys,N,N-dimethyl-

<220>

<221> SITE
 <222> (9)..(10)
 <223> Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 10 is Ser, Thr, Asn, Ala, Gly, Arg, Lys, ornithine, homoarginine, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-

<220>
 <221> SITE
 <222> (10)..(11)
 <223> Lys, any unnatural basic amino acid, His, halo-His, Pro or hydroxy-Pro; Xaa at residue 11 is Thr, Ser, Ala, Asp, Asn, Pro, hydroxy-Pro, Arg, ornithine, homoarginine, Lys, N-methyl-Lys,

<220>
 <221> SITE
 <222> (11)..(13)
 <223> N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 13 is Gly, Ser, Thr, Ala, Asn, Arg, ornithine, homoarginine, Lys, N-methyl-Lys,

<220>
 <221> SITE
 <222> (13)..(14)
 <223> N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 14 is Gln, Leu, His, halo-His, Trp (D or L), halo-Trp, neo-Trp, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-

<220>
 <221> SITE
 <222> (14)
 <223> Tyr, O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr, Arg, ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any unnatural basic amino acid or any unnatural aromatic amino

<220>
 <221> SITE
 <222> (14)..(15)
 <223> acid; Xaa at residue 15 is Asn, His, halo-His, Ile, Leu, Val, Gln, Arg, ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid;

<220>
 <221> SITE
 <222> (17)
 <223> Xaa at residue 17 is des-Xaa, Val, Ile, Leu, Arg, ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid.

<400> 1
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 1 5 10 15

Xaa

<210> 2

<211> 21
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<220>
 <223> Description of Artificial Sequence:Alpha-Conotoxin
 Peptide Generic Formula II.

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 <222> (1)..(3)
 <223> Xaa at residue 1 is des-Xaa, Asp, Glu or gamma-
 carboxy-Glu (Gla); Xaa at residue 2 is des-Xaa,
 Gln, Ala, Asp, Glu, Gla; Xaa at residue 3 is des-
 Xaa, Gly, Ala, Asp, Glu, Gla, Pro or hydroxy-Pro.

<220>
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 <222> (4)..(7)
 <223> Xaa at residue 4 is des-Xaa, Gly, Glu, Gla, Gln,
 Asp, Asn, Pro or hydroxy-Pro; Xaa at residue 7 is
 Ser, Thr, Gly, Glu, Gla, Asn, Trp (D or L),
 neo-Trp, halo-Trp, Arg, ornithine, homoarginine,

<220>
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 <222> (7)
 <223> Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-
 trimethyl-Lys, any unnatural basic amino acid,
 Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any

<220>
 <221> SITE
 <222> (7)..(8)
 <223> unnatural hydroxy containing amino acid; Xaa at
 residue 8 is Asp, Asn, His, halo-His, Thr, Ser,
 Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any

<220>
 <221> SITE
 <222> (8)..(10)
 <223> unnatural hydroxy containing amino acid; Xaa at
 residue 9 is Pro or hydroxy-Pro; Xaa at residue
 10 is Ala, Ser, Thr, Asp, Val, Ile, Pro, hydroxy-
 Pro, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,

<220>
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 <222> (10)..(12)
 <223> O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any
 unnatural hydroxy containing amino acid; Xaa at
 residue 12 is Gly, Ile, Leu, Val, Ala, Thr, Ser,
 Pro, hydroxy-Pro, Phe, Trp (D or L), neo-Trp,

<220>
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 <222> (12)..(13)
 <223> halo-Trp, Arg, ornithine, homoarginine, Lys, N-
 methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-
 Lys, any unnatural basic amino acid or any
 unnatural aromatic amino acid; Xaa at residue 13

<220>
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 <222> (13)
 <223> is Ala, Asn, Phe, Pro, hydroxy-Pro, Glu, Gla,
 Gln, His, halo-His, Val, Ser, Thr, Arg,
 ornithine, homoarginine, Lys, N-methyl-Lys, N,N-
 dimethyl-Lys, N,N,N-trimethyl-Lys or any

<220>
 <221> SITE
 <222> (13)..(14)
 <223> unnatural basic amino acid; Xaa at residue 14 is
 Thr, Ser, His, halo-His, Leu, Ile, Val, Asn, Met,
 Pro, hydroxy-Pro, Arg, ornithine, homoarginine,
 Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-

<220>
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 <222> (14)..(15)
 <223> trimethyl-Lys, any unnatural basic amino acid,
 Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any
 unnatural hydroxy containing amino acid; Xaa at

<220>
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 <222> (15)
 <223> residue 15 is Asn, Pro, hydroxy-Pro, Gln, Ser,
 Thr, Arg, ornithine, homoarginine, Lys, N-methyl-
 Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any
 unnatural basic amino acid, Tyr, nor-Tyr, mono-

<220>
 <221> SITE
 <222> (15)..(16)
 <223> halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-
 Tyr, nitro-Tyr or any unnatural hydroxy contain-
 ing amino acid; Xaa at residue 16 is des-Xaa,
 Gly, Thr, Ser, Pro, hydroxy-Pro, Tyr, nor-Tyr,

<220>
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 <222> (16)..(17)
 <223> mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy
 containing amino acid; Xaa at residue 17 is des-
 Xaa, Ile, Val, Asp, Leu, Phe, Arg, ornithine,

<220>
 <221> SITE
 <222> (17)
 <223> homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-
 Lys, N,N,N-trimethyl-Lys, any unnatural basic
 amino acid, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-
 Tyr, O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or

<220>
 <221> SITE
 <222> (17)..(19)
 <223> any unnatural hydroxy containing amino acid; Xaa
 at residue 19 is des-Xaa, Gly, Ala, Met, Ser,
 Thr, Trp (D or L), neo-Trp, halo-Trp, any
 unnatural aromatic amino acid, Arg, ornithine,

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<220>
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 <222> (19)..(20)
 <223> homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 20 is des-Xaa, Trp (D or L), neo-Trp, halo-Trp, any unnatural

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 <222> (20)..(21)
 <223> aromatic amino acid, Arg, ornithine, homo-arginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 21 is des-Xaa, Arg,

<220>
 <221> SITE
 <222> (21)
 <223> ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid.

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Xaa Cys Xaa Xaa Xaa
 20

<210> 3
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 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:Alpha-Conotoxin Peptide Generic Formula III.

<220>
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 <222> (1)..(3)
 <223> Xaa at residue 1 is des-Xaa, Ser or Thr; Xaa at residue 2 is des-Xaa, Asp, Glu, gamma-carboxy-Glu (Gla), Asn, Ser or Thr; Xaa at residue 3 is des-Xaa, Ala, Gly, Asn, Ser, Thr, Pro, hydroxy-Pro,

<220>
 <221> SITE
 <222> (3)..(4)
 <223> Arg, ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 4 is des-Xaa, Ala, Val, Leu, Ile, Gly, Glu, Gla, Gln,

<220>
 <221> SITE
 <222> (4)..(5)
 <223> Asp, Asn, Phe, Pro, hydroxy-Pro or any unnatural aromatic amino acid; Xaa at residue 5 is des-Xaa, Thr, Ser, Asp, Glu, Gla, Gln, Gly, Val, Asp, Asn, Ala, Pro, hydroxy-Pro, Arg, ornithine, homo-

<220>
 <221> SITE
 <222> (5)..(8)
 <223> arginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,
 N,N,N-trimethyl-Lys or any unnatural basic amino
 acid; Xaa at residue 8 is Thr, Ser, Asp, Asn, Met,
 Val, Ala, Gly, Leu, Ile, Phe, any unnatural

<220>
 <221> SITE
 <222> (8)..(9)
 <223> aromatic amino acid, Pro, hydroxy-Pro, Tyr, nor-
 Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy
 containing amino acid; Xaa at residue 9 is Ile,

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 <222> (9)
 <223> Leu, Val, Ser, Thr, Gln, Asn, Asp, Arg, His,
 halo-His, Phe, any unnatural aromatic amino
 acid, homoarginine, ornithine, Lys, N-methyl-Lys,
 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any

<220>
 <221> SITE
 <222> (9)..(10)
 <223> unnatural basic amino acid, Tyr, nor-Tyr, mono-
 halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-
 Tyr, nitro-Tyr or any unnatural hydroxy contain-
 ing amino acid; Xaa at residue 10 is Pro, hydroxy-

<220>
 <221> SITE
 <222> (10)
 <223> Pro, Ser, Thr, Ile, Asp, Leu, Val, Gly, Ala, Phe,
 any unnatural aromatic amino acid, Arg, ornithine,
 homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,
 N,N,N-trimethyl-Lys or any unnatural basic amino

<220>
 <221> SITE
 <222> (11)
 <223> acid; Xaa at residue 11 is Val, Ala, Gly, Ile,
 Leu, Asp, Ser, Thr, Pro, hydroxy-Pro, Arg,
 ornithine, homoarginine, Lys, N-methyl-Lys,
 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any

<220>
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 <222> (11)..(13)
 <223> unnatural basic amino acid; Xaa at residue 13 is
 His, halo-His, Arg, homoarginine, ornithine, Lys,
 N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-
 Lys, any unnatural basic amino acid, Asn, Ala,

<220>
 <221> SITE
 <222> (13)
 <223> Ser, Thr, Phe, Ile, Leu, Gly, Trp (D or L), neo-
 Trp, halo-Trp, any unnatural aromatic amino acid,
 Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any

<220>
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 <222> (13)..(14)
 <223> unnatural hydroxy containing amino acid; Xaa at residue 14 is Leu, Gln, Val, Ile, Gly, Met, Ala, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, Ser, Thr, Arg, homoarginine, ornithine, any unnatural basic amino acid, Asn, Glu, Ala, Gln, Phe, Trp (D or L), neo-Trp, halo-Trp or any unnatural aromatic amino acid; Xaa at residue 15 is Glu, Ala, Gln, Asn, Asp, Pro, hydroxy-Pro,

B29
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<220>
 <221> SITE
 <222> (15)
 <223> Ser, Gly, Thr, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, Arg, homoarginine, ornithine, any unnatural basic amino acid, Phe, His, halo-His, any unnatural aromatic acid, Leu,

<220>
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 <222> (15)..(16)
 <223> Met, Gly, Ala, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy containing amino acid; Xaa at residue 16 is His, halo-His, Asn, Thr,

<220>
 <221> SITE
 <222> (16)
 <223> Ser, Ile, Val, Leu, Phe, any unnatural aromatic amino acid, Arg, homoarginine, ornithine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any unnatural basic amino acid, Tyr, nor-Tyr,

<220>
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 <222> (16)..(17)
 <223> mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy containing amino acid; Xaa at residue 17 is Ser, Thr, Ala, Gln, Pro, hydroxy-Pro, Gly, Ile, Leu,

<220>
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 <222> (17)..(18)
 <223> Arg, ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural basic amino acid; Xaa at residue 18 is Asn, Glu, Ala, Asp, Gly, His, halo-His, Ala, Leu,

<220>
 <221> SITE
 <222> (18)
 <223> Gln, Arg, ornithine, homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any unnatural basic amino acid, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-

<220>
 <221> SITE
 <222> (18)..(19)
 <223> Tyr, nitro-Tyr or any unnatural hydroxy containing
 amino acid; Xaa at residue 19 is Met, Ile, Thr,
 Ser, Val, Leu, Pro, hydroxy-Pro, Phe, any
 unnatural aromatic amino acid, Tyr, nor-Tyr, mono-

<220>
 <221> SITE
 <222> (19)
 <223> halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-
 Tyr, nitro-Tyr, any unnatural hydroxy containing
 amino acid, Glu, Gla, Ala, His, halo-His, Arg,
 ornithine, homoarginine, Lys, N-methyl-Lys, N,N-

<220>
 <221> SITE
 <222> (21)
 <223> dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural
 basic amino acid; Xaa at residue 21 is des-Xaa,
 Gly, Asp, Asn, Ala, Ile, Leu, Ser, Thr, His, halo-
 His, Arg, ornithine, homoarginine, Lys, N-methyl-

<220>
 <221> SITE
 <222> (21)..(22)
 <223> Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any
 unnatural basic amino acid; Xaa at residue 22 is
 des-Xaa, Gly, Glu, Gla, Gln, Trp (D or L), neo-Trp,
 halo-Trp, any unnatural aromatic amino acid, Arg,

<220>
 <221> SITE
 <222> (22)..(23)
 <223> ornithine, homoarginine, Lys, N-methyl-Lys, N,N-
 dimethyl-Lys, N,N,N-trimethyl-Lys or any unnatural
 basic amino acid; Xaa at residue 23 is des-Xaa,
 Ser, Thr, Val, Ile, Ala, Arg, ornithine, homo-

<220>
 <221> SITE
 <222> (23)..(24)
 <223> arginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,
 N,N,N-trimethyl-Lys or any unnatural basic amino
 acid; Xaa at residue 24 is des-Xaa, Val, Asp, His,
 halo-His, Arg, ornithine, homoarginine, Lys, N-

<220>
 <221> SITE
 <222> (24)..(26)
 <223> methyl-Lys, N,N-dimethyl-Lys, N,N,N-trimethyl-Lys
 or any unnatural basic amino acid; Xaa at residue
 25 is des-Xaa, Asn, Pro or hydroxy-Pro; Xaa at
 residue 26 is des-Xaa, Arg, ornithine, homo-

<220>
 <221> SITE
 <222> (26)..(28)
 <223> arginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,
 N,N,N-trimethyl-Lys or any unnatural basic amino
 acid; Xaa at residue 27 is des-Xaa, Ser or Thr;
 Xaa at residue 28 is des-Xaa, Leu, Ile or Val.

<400> 3
 Xaa Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa
 1 5 10 15

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 20 25

<210> 4
 <211> 14
 <212> PRT
 <213> Conus imperialis

<220>
 <221> SITE
 <222> (2)..(11)
 <223> Xaa at residue 2 is Glu or gamma-carboxy-Glu; Xaa
 at residue 11 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 4
 Asp Xaa Cys Cys Ser Asp Ser Arg Cys Gly Xaa Asn Cys Leu
 1 5 10

<210> 5
 <211> 12
 <212> PRT
 <213> Conus imperialis

<220>
 <221> SITE
 <222> (10)
 <223> Xaa at residue 10 is Trp (D or L) or halo-Trp.

<400> 5
 Ala Cys Cys Ser Asp Arg Arg Cys Arg Xaa Arg Cys
 1 5 10

<210> 6
 <211> 13
 <212> PRT
 <213> Conus regius

<400> 6
 Phe Thr Cys Cys Arg Arg Gly Thr Cys Ser Gln His Cys
 1 5 10

<210> 7
 <211> 13
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (2)
 <223> Xaa at residue 2 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 7
 Asp Xaa Cys Cys Arg Arg His Ala Cys Thr Leu Ile Cys

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5

10

<210> 8
 <211> 13
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (2)..(8)
 <223> Xaa at residue 2 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr; Xaa at residues 7 and 8 is Pro or
 hydroxy-Pro.

<400> 8
 Asp Xaa Cys Cys Arg Arg Xaa Xaa Cys Thr Leu Ile Cys
 1 5 10

<210> 9
 <211> 13
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (6)..(10)
 <223> Xaa at residue 6 is Pro or hdroxy-Pro; Xaa at
 residue 10 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 9
 Gly Cys Cys Ser Asp Xaa Arg Cys Arg Xaa Arg Cys Arg
 1 5 10

<210> 10
 <211> 13
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (7)..(11)
 <223> Xaa at residue 7 is Pro or hydroxy-Pro; Xaa at
 residue 11 is Trp (D or L) or halo-Trp.

<400> 10
 Gly Gly Cys Cys Ser Asp Xaa Arg Cys Ala Xaa Arg Cys
 1 5 10

<210> 11
 <211> 17
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (3)..(10)
 <223> Xaa at residue 3 is Trp (D or L) or halo-Trp; Xaa

at residue 9 is Glu or gamma-carboxy-Glu; Xaa at residue 10 is Pro or hydroxy-Pro.

<220>
 <221> SITE
 <222> (15)
 <223> Xaa at residue 15 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 11
 Ile Ala Xaa Asp Ile Cys Cys Ser Xaa Xaa Asp Cys Asn His Xaa Cys
 1 5 10 15

Val

<210> 12
 <211> 12
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (6)..(9)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residue 9 is Lys, N-methyl-Lys, N,N-dimethyl-Lys
 or N,N,N-trimethyl-Lys.

<400> 12
 Gly Cys Cys Ser Asp Xaa Arg Cys Xaa His Gln Cys
 1 5 10

<210> 13
 <211> 14
 <212> PRT
 <213> Conus sponsalis

<220>
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 <222> (5)..(11)
 <223> Xaa at residues 5 and 11 is Pro or hydroxy-Pro;
 Xaa at residue 8 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 13
 Cys Cys Ser Asp Xaa Ala Cys Xaa Gln Thr Xaa Gly Cys Arg
 1 5 10

<210> 14
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 <212> PRT
 <213> Conus sponsalis

<220>
 <221> SITE
 <222> (3)..(5)
 <223> Xaa at residue 3 is Glu or gamma-carboxy-Glu; Xaa
 at residue 5 is Pro or hydroxy-Pro.

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 Cys Cys Xaa Asn Xaa Ala Cys Arg His Thr Gln Gly Cys

1

5

10

<210> 15
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 <212> PRT
 <213> Conus sulcatus

<220>
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 <222> (4)..(12)
 <223> Xaa at residue 4 is Trp or halo-Trp; Xaa at
 residue 6 is Pro or hydroxy-Pro; Xaa at residue 12
 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
 O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<400> 15
 Gly Cys Cys Xaa His Xaa Ala Cys Gly Arg His Xaa Cys
 1 5 10

<210> 16
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 <212> PRT
 <213> Conus achatinus

<220>
 <221> SITE
 <222> (2)..(11)
 <223> Xaa at residues 2 and 7 is Pro or hydroxy-Pro; Xaa
 at residue 11 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 16
 Ala Xaa Cys Cys Asn Asn Xaa Ala Cys Val Xaa His Arg Cys
 1 5 10

<210> 17
 <211> 15
 <212> PRT
 <213> Conus bullatus

<220>
 <221> SITE
 <222> (2)..(12)
 <223> Xaa at residues 2 and 8 is Pro or hydroxy-Pro; Xaa
 at residue 12 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 17
 Ala Xaa Gly Cys Cys Asn Asn Xaa Ala Cys Val Xaa His Arg Cys
 1 5 10 15

<210> 18
 <211> 14
 <212> PRT
 <213> Conus bullatus

<220>
 <221> SITE
 <222> (1)..(11)
 <223> Xaa at residues 1, 2 and 7 is Pro or hydroxy-Pro;

Xaa at residue 11 is Lys, N-methyl-Lys,
N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 18
Xaa Xaa Cys Cys Asn Asn Xaa Ala Cys Val Xaa His Arg Cys
1 5 10

<210> 19
<211> 16
<212> PRT
<213> Conus bullatus

<220>
<221> SITE
<222> (2)..(13)
<223> Xaa at residue 2 is Glu or gamma-carboxy-Glu; Xaa
at residue 6 is Trp or halo-Trp; Xaa at residues 8
11 and 13 is Pro or hydroxy-Pro.

<400> 19
Asp Xaa Asn Cys Cys Xaa Asn Xaa Ser Cys Xaa Arg Xaa Arg Cys Thr
1 5 10 15

<210> 20
<211> 13
<212> PRT
<213> Conus bullatus

<220>
<221> SITE
<222> (6)..(12)
<223> Xaa at residues 6 and 7 is Pro or hydroxy-Pro; Xaa
at residue 12 is Tyr, nor-Tyr, mono-halo-Tyr,
di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
nitro-Tyr.

<400> 20
Gly Cys Cys Ser Arg Xaa Xaa Cys Ala Val Leu Xaa Cys
1 5 10

<210> 21
<211> 13
<212> PRT
<213> Conus circumcissus

<220>
<221> SITE
<222> (6)
<223> Xaa at residue 6 is Pro or hydroxy-Pro.

<400> 21
Gly Cys Cys Gly Asn Xaa Asp Cys Thr Ser His Ser Cys
1 5 10

<210> 22
<211> 16
<212> PRT
<213> Conus stercusmuscarum

<220>

<221> SITE
 <222> (6)..(11)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residue 11 is Glu or gamma-carboxy-Glu.

<400> 22
 Gly Cys Cys Ser Asn Xaa Val Cys His Leu Xaa His Ser Asn Met Cys
 1 5 10 15

<210> 23
 <211> 17
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residue 14 is Glu or gamma-carboxy-Glu; Xaa at
 residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>
 <221> SITE
 <222> (15)
 <223> nitro-Tyr.

<400> 23
 Gly Cys Cys Ser Asn Xaa Val Cys Arg Gln Asn Asn Ala Xaa Xaa Cys
 1 5 10 15

Arg

<210> 24
 <211> 18
 <212> PRT
 <213> Conus textile

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 24
 Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile
 1 5 10 15

Cys Arg

<210> 25
 <211> 18
 <212> PRT
 <213> Conus radiatus

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;

Xaa at residues 2 and 15 is Glu or
gamma-carboxy-Glu.

<400> 25

Xaa Xaa Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile
1 5 10 15

Cys Arg

<210> 26

<211> 18

<212> PRT

<213> Conus radiatus

<220>

<221> SITE

<222> (1)..(15)

<223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 26

Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile
1 5 10 15

Cys Asp

<210> 27

<211> 18

<212> PRT

<213> Conus omaria

<220>

<221> SITE

<222> (1)..(15)

<223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 27

Xaa Arg Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile
1 5 10 15

Cys Arg

<210> 28

<211> 18

<212> PRT

<213> Conus omaria

<220>

<221> SITE

<222> (1)..(14)

<223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro.

<400> 28

Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Gly Ile
1 5 10 15

Cys Arg

<210> 29
 <211> 18
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 29
 Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Thr
 1 5 10 15

Cys Arg

<210> 30
 <211> 18
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 30
 Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Val
 1 5 10 15

Cys Arg

<210> 31
 <211> 18
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 31
 Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Ile Asp His Xaa Xaa Ile
 1 5 10 15

Cys Arg

<210> 32
 <211> 21
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 32
 Xaa Gln Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile
 1 5 10 15
 Cys Arg Arg Arg Arg
 20

<210> 33
 <211> 17
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (7)..(15)
 <223> Xaa at residues 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 33
 Gly Gly Cys Cys Ser His Xaa Ala Cys Ala Val Asn His Xaa Xaa Leu
 1 5 10 15
 Cys

<210> 34
 <211> 16
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 14 is Glu or gamma-carboxy-Glu.

<400> 34
 Gly Cys Cys Ser His Xaa Ala Cys Ser Val Asn His Xaa Xaa Leu Cys
 1 5 10 15

<210> 35
 <211> 16
 <212> PRT
 <213> Conus dalli

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 14 is Glu or gamma-carboxy-Glu.

<400> 35
 Gly Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile Cys
 1 5 10 15

<210> 36
 <211> 19
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (6)..(18)
 <223> Xaa at residues 6 and 15 is Pro or hydroxy-Pro;
 Xaa at residue 11 is Lys, N,-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys; Xaa at
 residues 14 and 18 is Glu or gamma-carboxy-Glu.

<400> 36
 Gly Cys Cys Ser His Xaa Ala Cys Ser Gly Xaa Thr Gln Xaa Xaa Cys
 1 5 10 15

Arg Xaa Ser

<210> 37
 <211> 18
 <212> PRT
 <213> Conus tulipa

<220>
 <221> SITE
 <222> (1)..(14)
 <223> Xaa at residues 1, 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 14 is Glu or gamma-carboxy-Glu.

<400> 37
 Xaa Cys Cys Ser His Xaa Ala Cys Ser Gly Asn Asn Xaa Xaa Phe Cys
 1 5 10 15

Arg Gln

<210> 38
 <211> 18
 <212> PRT
 <213> Conus tulipa

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 14 is Glu or gamma-carboxy-Glu.

<400> 38
 Gly Cys Cys Ser His Xaa Ala Cys Ser Gly Asn Asn Xaa Xaa Phe Cys
 1 5 10 15

Arg Gln

<210> 39
 <211> 16
 <212> PRT
 <213> Conus pennaceus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 39
 Gly Cys Cys Ser His Xaa Xaa Cys Ala Met Asn Asn Xaa Asp Xaa Cys
 1 5 10 15

<210> 40
 <211> 16
 <212> PRT
 <213> Conus pennaceus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residuew 6, 7 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 40
 Gly Cys Cys Ser His Xaa Xaa Cys Phe Leu Asn Asn Xaa Asp Xaa Cys
 1 5 10 15

<210> 41
 <211> 17
 <212> PRT
 <213> Conus textile

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 11 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N.N.N-trimethyl-Lys.

<400> 41
 Gly Cys Cys Ser Asn Xaa Xaa Cys Ile Ala Xaa Asn Xaa His Met Cys
 1 5 10 15

Gly

<210> 42
 <211> 16
 <212> PRT
 <213> Conus distans

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro.

<400> 42
 Gly Cys Cys Ser Asn Xaa Xaa Cys Ala His Asn Asn Xaa Asp Cys Arg
 1 5 10 15

<210> 43
 <211> 17
 <212> PRT
 <213> Conus tulipa

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 43
 Gly Cys Cys Ser Asn Xaa Ala Cys Ala Gly Asn Asn Xaa His Val Cys
 1 5 10 15

Arg

<210> 44
 <211> 16
 <212> PRT
 <213> Conus dalli

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 44
 Gly Cys Cys Ser Arg Xaa Ala Cys Ile Ala Asn Asn Xaa Asp Leu Cys
 1 5 10 15

<210> 45
 <211> 20
 <212> PRT
 <213> Conus circumciscus

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residues 11 and 14 is Glu or
 gamma-carboxy-Glu.

<400> 45
 Gly Cys Cys Ser Asn Xaa Val Cys His Val Xaa His Xaa Xaa Leu Cys
 1 5 10 15

Arg Arg Arg Arg
 20

<210> 46
 <211> 18
 <212> PRT
 <213> Conus sulcatus

<220>
 <221> SITE
 <222> (7)..(15)
 <223> Xaa at residues 7, 12 and 14 is Pro or
 hydroxy-Pro; Xaa at residue 11 is Lys,

N-methyl-Lys, N,N-dimethyl-Lys or
N,N,N-trimethyl-Lys; Xaa at residue 15 is Glu or

<220>
<221> SITE
<222> (7)..(15)
<223> gamma-carboxy-Glu.

<400> 46
Gly Gly Cys Cys Ser Phe Xaa Ala Cys Arg Xaa Xaa Arg Xaa Xaa Met
1 5 10 15

Cys Gly

<210> 47
<211> 18
<212> PRT
<213> Conus textile

<220>
<221> SITE
<222> (1)..(15)
<223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro;
Xaa at residues 2 and 15 is Glu or
gamma-carboxy-Glu.

<400> 47
Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Asn Ser Ser His Xaa Xaa Leu
1 5 10 15

Cys Arg

<210> 48
<211> 18
<212> PRT
<213> Conus dalli

<220>
<221> SITE
<222> (1)..(15)
<223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-pro;
Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 48
Xaa Gln Cys Cys Ser Asp Xaa Arg Cys Asn Val Gly His Xaa Xaa Leu
1 5 10 15

Cys Gly

<210> 49
<211> 18
<212> PRT
<213> Conus dalli

<220>
<221> SITE
<222> (1)..(15)
<223> Xaa at residue 1 is Gln or pyro-Glu; Xaa at

residues 7 and 14 is Pro or hydroxy-Pro; Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 49

Xaa Val Cys Cys Ser Asp Xaa Arg Cys Asn Val Gly His Xaa Xaa Ile
1 5 10 15

Cys Gly

<210> 50

<211> 16

<212> PRT

<213> Conus textile

<220>

<221> SITE

<222> (6)..(13)

<223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro.

<400> 50

Gly Cys Cys Ser Arg Xaa Xaa Cys Ile Ala Asn Asn Xaa Asp Leu Cys
1 5 10 15

<210> 51

<211> 18

<212> PRT

<213> Conus omaria

<220>

<221> SITE

<222> (1)..(15)

<223> Xaa at residues 1 and 14 is Pro or hydroxy-Pro;
Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 51

Xaa Gln Cys Cys Ser His Leu Ala Cys Asn Val Asp His Xaa Xaa Ile
1 5 10 15

Cys Arg

<210> 52

<211> 19

<212> PRT

<213> Conus sulcatus

<220>

<221> SITE

<222> (5)..(14)

<223> Xaa at residue 5 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr; Xaa at residue 13 is Pro or hydroxy-Pro; Xaa at residue 14 is Glu or

<220>

<221> SITE

<222> (14)..(18)

<223> gamma-carboxy-Glu; Xaa at residue 18 is Trp or halo-Trp.

<400> 52

Gly Cys Cys Ser Xaa Phe Asp Cys Arg Met Met Phe Xaa Xaa Met Cys
 1 5 10 15

Gly Xaa Arg

<210> 53

<211> 18

<212> PRT

<213> Conus sulcatus

<220>

<221> SITE

<222> (11)..(12)

<223> Xaa at residue 11 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys; Xaa at
 residue 12 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>

<221> SITE

<222> (12)..(15)

<223> nitro-Tyr; Xaa at residue 14 is Pro or hydroxy-
 Pro; Xaa at residue 15 is Glu or gamma-carboxy-
 Glu.

<400> 53

Gly Gly Cys Cys Ser Phe Ala Ala Cys Arg Xaa Xaa Arg Xaa Xaa Met
 1 5 10 15

Cys Gly

<210> 54

<211> 20

<212> PRT

<213> Conus sulcatus

<220>

<221> SITE

<222> (7)..(15)

<223> Xaa at residue 7 is Pro or hydroxy-Pro; Xaa at
 residue 10 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr; Xaa at residue 15 is Glu or

<220>

<221> SITE

<222> (7)..(15)

<223> gamma-carboxy-Glu.

<400> 54

Gly Gly Cys Cys Phe His Xaa Val Cys Xaa Ile Asn Leu Leu Xaa Met
 1 5 10 15

Cys Arg Gln Arg
 20

<210> 55

<211> 19

<212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (7)..(15)
 <223> Xaa at residues 7, 11 and 14 is Tyr, nor-Tyr,
 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr; Xaa at residues 8, 9 and 15 is Pro
 or hydroxy-Pro.

<220>
 <221> SITE
 <222> (12)..(16)
 <223> Xaa at residues 12 and 16 is Glu or
 gamma-carboxy-Glu.

<400> 55
 Ser Ala Thr Cys Cys Asn Xaa Xaa Xaa Cys Xaa Xaa Thr Xaa Xaa Xaa
 1 5 10 15

Ser Cys Leu

<210> 56
 <211> 17
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (5)..(13)
 <223> Xaa at residues 5 and 12 is Tyr, no-Tyr,
 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr or nitro-Tyr; Xaa at residues 6, 7
 and 13 is Pro or hydroxy-Pro.

<220>
 <221> SITE
 <222> (10)..(14)
 <223> Xaa at residues 10 and 14 is Glu or
 gamma-carboxy-Glu.

<400> 56
 Ala Cys Cys Ala Xaa Xaa Xaa Cys Phe Xaa Ala Xaa Xaa Xaa Arg Cys
 1 5 10 15

Leu

<210> 57
 <211> 19
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (3)..(16)
 <223> Xaa at residues 3, 12 and 16 is Glu or
 gamma-carboxy-Glu; Xaa at residues 6, 7, 11 and 14
 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
 O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<220>
 <221> SITE
 <222> (8)..(15)
 <223> Xaa at residues 8, 9 and 15 is Pro or hydroxy-Pro.

<400> 57
 Asn Ala Xaa Cys Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Ala Xaa Xaa Xaa
 1 5 10 15

Ile Cys Leu

<210> 58
 <211> 227
 <212> DNA
 <213> Conus magus

<220>
 <221> CDS
 <222> (1)..(189)

<400> 58
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

 ttc cct tca gat cgt gca tct gat ggc agg aat gcc gca gcc aac gac 96
 Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp
 20 25 30

 aaa gcg tct gac gtg atc acg ctg gcc ctc aag gga tgc tgt tcc aac 144
 Lys Ala Ser Asp Val Ile Thr Leu Ala Leu Lys Gly Cys Cys Ser Asn
 35 40 45

 cct gtc tgt cac ttg gag cat tca aac ctt tgt ggt aga aga cgc 189
 Pro Val Cys His Leu Glu His Ser Asn Leu Cys Gly Arg Arg Arg
 50 55 60

 tgatgctcca ggaccctctg aaccacgacg ttcgagca 227

<210> 59
 <211> 63
 <212> PRT
 <213> Conus magus

<400> 59
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

 Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp
 20 25 30

 Lys Ala Ser Asp Val Ile Thr Leu Ala Leu Lys Gly Cys Cys Ser Asn
 35 40 45

 Pro Val Cys His Leu Glu His Ser Asn Leu Cys Gly Arg Arg Arg
 50 55 60

<210> 60
 <211> 208
 <212> DNA

<213> Conus aulicus

<220>

<221> CDS

<222> (1)..(168)

<400> 60

atg	ttc	acc	gtg	ttt	ctg	ttg	gtt	gtc	ttg	gca	acc	acc	gtc	gtt	tcc	48
Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5					10					15		

ttc	act	tca	gat	cgt	gca	tct	gat	ggc	agg	aag	gac	gca	gcg	tct	ggc	96
Phe	Thr	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Lys	Asp	Ala	Ala	Ser	Gly	
			20					25					30			

ctg	atc	gct	ctg	acc	atc	aag	gga	tgc	tgt	tct	tat	cct	ccc	tgt	ttc	144
Leu	Ile	Ala	Leu	Thr	Ile	Lys	Gly	Cys	Cys	Ser	Tyr	Pro	Pro	Cys	Phe	
		35					40					45				

gcg	act	aat	tca	gac	tat	tgt	ggc	tgacgacgct	gatgctccag	gaccctctga	198
Ala	Thr	Asn	Ser	Asp	Tyr	Cys	Gly				
	50					55					

accacgacgt	208
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<210> 61

<211> 56

<212> PRT

<213> Conus aulicus

<400> 61

Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser
1				5					10					15	

Phe	Thr	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Lys	Asp	Ala	Ala	Ser	Gly
			20					25					30		

Leu	Ile	Ala	Leu	Thr	Ile	Lys	Gly	Cys	Cys	Ser	Tyr	Pro	Pro	Cys	Phe
		35					40					45			

Ala	Thr	Asn	Ser	Asp	Tyr	Cys	Gly
	50					55	

<210> 62

<211> 205

<212> DNA

<213> Conus aulicus

<220>

<221> CDS

<222> (1)..(174)

<400> 62

atg	ttc	acc	gtg	ttt	ctg	ttg	gtc	gtc	ttg	gca	acc	acc	gtc	gtt	tcc	48
Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5					10					15		

ttc	act	tca	gat	cgt	gca	tct	gat	ggc	agg	aag	gac	gca	gcg	tct	ggc	96
Phe	Thr	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Lys	Asp	Ala	Ala	Ser	Gly	
			20					25					30			

ctg	att	gct	ctg	acc	atg	aag	gga	tgc	tgt	tct	tat	cct	ccc	tgt	ttc	144
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

Leu Ile Ala Leu Thr Met Lys Gly Cys Cys Ser Tyr Pro Pro Cys Phe
 35 40 45

gcg act aat cca gac tgt ggt cga cga cgc tgatgctcca ggaccctctg 194
 Ala Thr Asn Pro Asp Cys Gly Arg Arg Arg
 50 55

aaccacgacg t 205

<210> 63
 <211> 58
 <212> PRT
 <213> Conus aulicus

<400> 63
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Phe Thr Ser Asp Arg Ala Ser Asp Gly Arg Lys Asp Ala Ala Ser Gly
 20 25 30

Leu Ile Ala Leu Thr Met Lys Gly Cys Cys Ser Tyr Pro Pro Cys Phe
 35 40 45

Ala Thr Asn Pro Asp Cys Gly Arg Arg Arg
 50 55

<210> 64
 <211> 223
 <212> DNA
 <213> Conus textile

<220>
 <221> CDS
 <222> (1)..(192)

<400> 64
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ttc tct tca ggt cgt agt aca ttt cgt ggc agg aat gcc gca gcc aaa 96
 Phe Ser Ser Gly Arg Ser Thr Phe Arg Gly Arg Asn Ala Ala Ala Lys
 20 25 30

gcg tct ggc ctg gtc agt ctg act gac agg aga cca gaa tgc tgt agt 144
 Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Glu Cys Cys Ser
 35 40 45

gat cct cgc tgt aac tcg agt cat cca gaa ctt tgt ggt gga aga cgc 192
 Asp Pro Arg Cys Asn Ser Ser His Pro Glu Leu Cys Gly Gly Arg Arg
 50 55 60

tgatgctcca ggaccctctg aaccacgacg t 223

<210> 65
 <211> 64
 <212> PRT
 <213> Conus textile

<400> 65

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Ser Ser Gly Arg Ser Thr Phe Arg Gly Arg Asn Ala Ala Ala Lys
20 25 30

Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Glu Cys Cys Ser
35 40 45

Asp Pro Arg Cys Asn Ser Ser His Pro Glu Leu Cys Gly Gly Arg Arg
50 55 60

<210> 66

<211> 244

<212> DNA

<213> Conus textile

<220>

<221> CDS

<222> (1)..(168)

<400> 66

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc gcc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Ala Val Val Ser
1 5 10 15

ttc act tca gat cgt gca tct gat gac ggg aaa gcc gct gcg tct gac 96
Phe Thr Ser Asp Arg Ala Ser Asp Asp Gly Lys Ala Ala Ala Ser Asp
20 25 30

ctg atc act ctg acc atc aag gga tgc tgt tct cgt cct ccc tgt atc 144
Leu Ile Thr Leu Thr Ile Lys Gly Cys Cys Ser Arg Pro Pro Cys Ile
35 40 45

gcg aat aat cca gac ttg tgt ggt tgacgacgct gatgctccag aacggtctga 198
Ala Asn Asn Pro Asp Leu Cys Gly
50 55

accacgacgt tcgagcaatg ttcaccgtgt ttctgttggt tgtctt 244

<210> 67

<211> 56

<212> PRT

<213> Conus textile

<400> 67

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Ala Val Val Ser
1 5 10 15

Phe Thr Ser Asp Arg Ala Ser Asp Asp Gly Lys Ala Ala Ala Ser Asp
20 25 30

Leu Ile Thr Leu Thr Ile Lys Gly Cys Cys Ser Arg Pro Pro Cys Ile
35 40 45

Ala Asn Asn Pro Asp Leu Cys Gly
50 55

<210> 68

<211> 223

<212> DNA

<213> Conus textile

<220>

<221> CDS

<222> (1)..(183)

<400> 68

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atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
  1             5             10             15

ttc act tca ggt cgt agt aca ttt cgt ggc agg aat gcc gca gcc aaa 96
Phe Thr Ser Gly Arg Ser Thr Phe Arg Gly Arg Asn Ala Ala Ala Lys
             20             25             30

gcg tct ggc ctg gtc agt ctg act gac agg aga cca caa tgc tgt tct 144
Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Gln Cys Cys Ser
             35             40             45

cat cct gcc tgt aac gta gat cat cca gaa att tgt cgt tgaagacgct 193
His Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Arg
      50             55             60

gatgctccag gaccctctga accacgacgt 223

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<210> 69

<211> 61

<212> PRT

<213> Conus textile

<400> 69

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Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
  1             5             10             15

Phe Thr Ser Gly Arg Ser Thr Phe Arg Gly Arg Asn Ala Ala Ala Lys
             20             25             30

Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Gln Cys Cys Ser
             35             40             45

His Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Arg
      50             55             60

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<210> 70

<211> 223

<212> DNA

<213> Conus radiatus

<220>

<221> CDS

<222> (1)..(183)

<400> 70

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atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
  1             5             10             15

ttc act tca ggt cgt cgt aca ttt cat ggc agg aat gcc gca gcc aaa 96
Phe Thr Ser Gly Arg Arg Thr Phe His Gly Arg Asn Ala Ala Ala Lys
             20             25             30

gcg tct ggc ctg gtc agt ctg act gac agg aga cca gaa tgc tgt tct 144

```

Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Glu Cys Cys Ser
 35 40 45

cat cct gcc tgt aac gta gat cat cca gaa att tgt cgt tgaagacgct 193
 His Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Arg
 50 55 60

gatgctccag gaccctctga accacgacgt 223

<210> 71
 <211> 61
 <212> PRT
 <213> Conus radiatus

<400> 71
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Phe Thr Ser Gly Arg Arg Thr Phe His Gly Arg Asn Ala Ala Ala Lys
 20 25 30

Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Glu Cys Cys Ser
 35 40 45

His Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Arg
 50 55 60

<210> 72
 <211> 223
 <212> DNA
 <213> Conus radiatus

<220>
 <221> CDS
 <222> (1)..(183)

<400> 72
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ttc act tca ggt cgt agt aca ttt cgt ggc agg aat gcc gca gcc aaa 96
 Phe Thr Ser Gly Arg Ser Thr Phe Arg Gly Arg Asn Ala Ala Ala Lys
 20 25 30

gcg tct ggc ctg gtc agt ctg act gac agg aga cca caa tgc tgt tct 144
 Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Gln Cys Cys Ser
 35 40 45

cat cct gcc tgt aac gta gat cat cca gaa att tgc gat tgaagacgct 193
 His Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Asp
 50 55 60

gatgctccag gaccctctga accacgacgt 223

<210> 73
 <211> 61
 <212> PRT
 <213> Conus radiatus

<400> 73

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Thr Ser Gly Arg Ser Thr Phe Arg Gly Arg Asn Ala Ala Ala Lys
20 25 30

Ala Ser Gly Leu Val Ser Leu Thr Asp Arg Arg Pro Gln Cys Cys Ser
35 40 45

His Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Asp
50 55 60

<210> 74

<211> 218

<212> DNA

<213> Conus striatus

<220>

<221> CDS

<222> (1)..(171)

<400> 74

atg ttc act gtg ttt ctg ttg gtt gtc ttg gca atc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Ile Thr Val Val Ser
1 5 10 15

ttc cct tta gat cgt gaa tct gat ggc gcg aat gcc gaa gcc cgc acc 96
Phe Pro Leu Asp Arg Glu Ser Asp Gly Ala Asn Ala Glu Ala Arg Thr
20 25 30

cac gat cat gag aag cac gca ctg gac cgg aat gga tgc tgt agg aat 144
His Asp His Glu Lys His Ala Leu Asp Arg Asn Gly Cys Cys Arg Asn
35 40 45

cct gcc tgt gag agc cac aga tgt ggt tgacgacgct gatgctccag 191
Pro Ala Cys Glu Ser His Arg Cys Gly
50 55

gaccctctga accacgacgt tcgagca 218

<210> 75

<211> 57

<212> PRT

<213> Conus striatus

<400> 75

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Ile Thr Val Val Ser
1 5 10 15

Phe Pro Leu Asp Arg Glu Ser Asp Gly Ala Asn Ala Glu Ala Arg Thr
20 25 30

His Asp His Glu Lys His Ala Leu Asp Arg Asn Gly Cys Cys Arg Asn
35 40 45

Pro Ala Cys Glu Ser His Arg Cys Gly
50 55

<210> 76

<211> 227

<212> DNA

<213> Conus bandanus

<220>

<221> CDS

<222> (1)..(180)

<400> 76

atg	ttc	acc	atg	ttt	ctg	ttg	gtt	gtc	ttg	gca	acc	act	gtc	gtt	tcc	48
Met	Phe	Thr	Met	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5				10					15			

ttc	gct	tca	gat	cgt	gca	tct	gat	ggc	agg	aat	gcc	gca	gcc	aag	gac	96
Phe	Ala	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Lys	Asp	
			20					25					30			

aaa	gcg	tct	gac	ctg	gtc	gct	ctg	acc	gtc	aag	gga	tgc	tgt	tct	cat	144
Lys	Ala	Ser	Asp	Leu	Val	Ala	Leu	Thr	Val	Lys	Gly	Cys	Cys	Ser	His	
		35				40						45				

cct	gcc	tgt	agc	gtg	aat	aat	cca	gac	att	tgt	ggc	tgaagacgct	190
Pro	Ala	Cys	Ser	Val	Asn	Asn	Pro	Asp	Ile	Cys	Gly		
	50					55					60		

gatgctccag	gacctctga	accacgacgt	tcgagca	227
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<210> 77

<211> 60

<212> PRT

<213> Conus bandanus

<400> 77

Met	Phe	Thr	Met	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser
1				5					10					15	

Phe	Ala	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Lys	Asp
			20					25					30		

Lys	Ala	Ser	Asp	Leu	Val	Ala	Leu	Thr	Val	Lys	Gly	Cys	Cys	Ser	His
		35				40						45			

Pro	Ala	Cys	Ser	Val	Asn	Asn	Pro	Asp	Ile	Cys	Gly
	50					55					60

<210> 78

<211> 104

<212> DNA

<213> Conus bandanus

<220>

<221> CDS

<222> (1)..(54)

<400> 78

aaa	gaa	tgc	tgt	act	cat	cct	gcc	tgt	cac	gtg	agt	cat	cca	gaa	ctc	48
Lys	Glu	Cys	Cys	Thr	His	Pro	Ala	Cys	His	Val	Ser	His	Pro	Glu	Leu	
1				5					10					15		

tgt	ggc	tga	aaagcga	cgtgacgctc	caggaccctc	tgaaccacga	cgttcgagca	104
Cys	Gly							

<210> 79

<211> 18
 <212> PRT
 <213> Conus bandanus

<400> 79
 Lys Glu Cys Cys Thr His Pro Ala Cys His Val Ser His Pro Glu Leu
 1 5 10 15
 Cys Gly

<210> 80
 <211> 206
 <212> DNA
 <213> Conus bandanus

<220>
 <221> CDS
 <222> (1)..(171)

<400> 80
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca act gct gtt ctt cca 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Ala Val Leu Pro
 1 5 10 15
 gtc act tta gat cgt gca tct gat gga agg aat gca gca gcc aac gcc 96
 Val Thr Leu Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala
 20 25 30
 aaa acg cct cgc ctg atc gcg cca ttc atc agg gat tat tgc tgt cat 144
 Lys Thr Pro Arg Leu Ile Ala Pro Phe Ile Arg Asp Tyr Cys Cys His
 35 40 45
 aga ggt ccc tgt atg gta tgg tgt ggt tgaagccgct gctgctccag 191
 Arg Gly Pro Cys Met Val Trp Cys Gly
 50 55
 gaccctctga accac 206

<210> 81
 <211> 57
 <212> PRT
 <213> Conus bandanus

<400> 81
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Ala Val Leu Pro
 1 5 10 15
 Val Thr Leu Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala
 20 25 30
 Lys Thr Pro Arg Leu Ile Ala Pro Phe Ile Arg Asp Tyr Cys Cys His
 35 40 45
 Arg Gly Pro Cys Met Val Trp Cys Gly
 50 55

<210> 82
 <211> 174
 <212> DNA
 <213> Conus characteristicus

<220>
 <221> CDS
 <222> (1)..(171)

<400> 82

atg	ttc	acc	gtg	ttt	ctg	ttg	gtt	gtc	ttg	gca	acc	act	gtg	gtt	tcc	48
Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5				10					15			
ttc	act	tca	gat	cgt	gct	tct	gat	ggc	agg	aat	gcc	gca	gcc	aac	gcg	96
Phe	Thr	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Asn	Ala	
			20					25					30			
ttt	gac	ctg	atc	gct	ctg	atc	gcc	agg	caa	aat	tgc	tgt	agc	att	ccc	144
Phe	Asp	Leu	Ile	Ala	Leu	Ile	Ala	Arg	Gln	Asn	Cys	Cys	Ser	Ile	Pro	
		35					40					45				
agc	tgt	tgg	gag	aaa	tat	aaa	tgt	agt	taa							174
Ser	Cys	Trp	Glu	Lys	Tyr	Lys	Cys	Ser								
	50					55										

<210> 83
 <211> 57
 <212> PRT
 <213> Conus characteristicus

<400> 83

Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5				10					15			
Phe	Thr	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Asn	Ala	
			20					25					30			
Phe	Asp	Leu	Ile	Ala	Leu	Ile	Ala	Arg	Gln	Asn	Cys	Cys	Ser	Ile	Pro	
		35					40					45				
Ser	Cys	Trp	Glu	Lys	Tyr	Lys	Cys	Ser								
	50					55										

<210> 84
 <211> 219
 <212> DNA
 <213> Conus characteristicus

<220>
 <221> CDS
 <222> (1)..(189)

<400> 84

atg	ttc	acc	gtg	ttt	ctg	ttg	gtt	gtc	ttg	gca	acc	act	gtg	gtt	tcc	48
Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5				10					15			
ttc	act	tca	gat	cgt	gcg	tct	gaa	ggc	agg	aat	gct	gca	gcc	aag	gac	96
Phe	Thr	Ser	Asp	Arg	Ala	Ser	Glu	Gly	Arg	Asn	Ala	Ala	Ala	Lys	Asp	
			20					25					30			
aaa	gcg	tct	gac	ctg	gtg	gct	ctg	aca	gtc	agg	gga	tgc	tgt	gcc	att	144
Lys	Ala	Ser	Asp	Leu	Val	Ala	Leu	Thr	Val	Arg	Gly	Cys	Cys	Ala	Ile	
		35					40					45				
cgt	gaa	tgt	cgc	ttg	cag	aat	gca	gcg	tat	tgt	ggt	gga	ata	tac		189

Arg Glu Cys Arg Leu Gln Asn Ala Ala Tyr Cys Gly Gly Ile Tyr
 50 55 60

tgatgctcca ggaccctctg aaccacgacg

219

<210> 85
 <211> 63
 <212> PRT
 <213> Conus characteristicus

<400> 85
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Thr Ser Asp Arg Ala Ser Glu Gly Arg Asn Ala Ala Ala Lys Asp
 20 25 30
 Lys Ala Ser Asp Leu Val Ala Leu Thr Val Arg Gly Cys Cys Ala Ile
 35 40 45
 Arg Glu Cys Arg Leu Gln Asn Ala Ala Tyr Cys Gly Gly Ile Tyr
 50 55 60

<210> 86
 <211> 217
 <212> DNA
 <213> Conus tulipa

<220>
 <221> CDS
 <222> (1)..(186)

<400> 86
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc cct tca gat att gca act gag ggc agg aat gcc gca gcc aaa gcg 96
 Phe Pro Ser Asp Ile Ala Thr Glu Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 ttt gac ctg ata tct tcg atc gtc aag aaa gga tgc tgt tcc cat cct 144
 Phe Asp Leu Ile Ser Ser Ile Val Lys Lys Gly Cys Cys Ser His Pro
 35 40 45
 gcc tgt tcg ggg aat aat cca gaa ttt tgt cgt caa ggt cgc 186
 Ala Cys Ser Gly Asn Asn Pro Glu Phe Cys Arg Gln Gly Arg
 50 55 60

tgatgctcca ggaccctctg aaccacgacg t

217

<210> 87
 <211> 62
 <212> PRT
 <213> Conus tulipa

<400> 87
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Ile Ala Thr Glu Gly Arg Asn Ala Ala Ala Lys Ala

20 25 30
 Phe Asp Leu Ile Ser Ser Ile Val Lys Lys Gly Cys Cys Ser His Pro
 35 40 45

Ala Cys Ser Gly Asn Asn Pro Glu Phe Cys Arg Gln Gly Arg
 50 55 60

<210> 88
 <211> 217
 <212> DNA
 <213> Conus tulipa

<220>
 <221> CDS
 <222> (1)..(186)

<400> 88
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc cct tca gat ata gca act gag ggc agg aat gcc gca gcc aaa gcg 96
 Phe Pro Ser Asp Ile Ala Thr Glu Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 ttt gac ctg ata tct tcg atc gtc agg aaa gga tgc tgt tcc aat ccc 144
 Phe Asp Leu Ile Ser Ser Ile Val Arg Lys Gly Cys Cys Ser Asn Pro
 35 40 45
 gcc tgt gcg ggg aat aat cca cat gtt tgt cgt caa ggt cgc 186
 Ala Cys Ala Gly Asn Asn Pro His Val Cys Arg Gln Gly Arg
 50 55 60
 tgatgctcca ggaccctctg aaccacgacg t 217

<210> 89
 <211> 62
 <212> PRT
 <213> Conus tulipa

<400> 89
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Ile Ala Thr Glu Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 Phe Asp Leu Ile Ser Ser Ile Val Arg Lys Gly Cys Cys Ser Asn Pro
 35 40 45
 Ala Cys Ala Gly Asn Asn Pro His Val Cys Arg Gln Gly Arg
 50 55 60

<210> 90
 <211> 226
 <212> DNA
 <213> Conus sulcatus

<220>
 <221> CDS

<222> (1)..(195)

<400> 90

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atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
  1             5             10             15

ttc aat tca gat cgt gat cca gca tta ggt ggc agg aat gct gca gcc 96
Phe Asn Ser Asp Arg Asp Pro Ala Leu Gly Gly Arg Asn Ala Ala Ala
          20             25             30

aaa gcg tct gac aag atc gct tcg acc ctc aag aga aga gga tgc tgt 144
Lys Ala Ser Asp Lys Ile Ala Ser Thr Leu Lys Arg Arg Gly Cys Cys
          35             40             45

tcg tat ttt gac tgt aga atg atg ttt cca gaa atg tgt ggt tgg cga 192
Ser Tyr Phe Asp Cys Arg Met Met Phe Pro Glu Met Cys Gly Trp Arg
          50             55             60

ggc tgatgctcca ggaccctctg aaccacgacg t 226
Gly
65

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<210> 91

<211> 65

<212> PRT

<213> Conus sulcatus

<400> 91

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Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
  1             5             10             15

Phe Asn Ser Asp Arg Asp Pro Ala Leu Gly Gly Arg Asn Ala Ala Ala
          20             25             30

Lys Ala Ser Asp Lys Ile Ala Ser Thr Leu Lys Arg Arg Gly Cys Cys
          35             40             45

Ser Tyr Phe Asp Cys Arg Met Met Phe Pro Glu Met Cys Gly Trp Arg
          50             55             60

Gly
65

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<210> 92

<211> 226

<212> DNA

<213> Conus sulcatus

<220>

<221> CDS

<222> (1)..(195)

<400> 92

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atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
  1             5             10             15

ttc aat tca gat cgt gat cca gca tta ggt ggc agg aat gct gca gcc 96
Phe Asn Ser Asp Arg Asp Pro Ala Leu Gly Gly Arg Asn Ala Ala Ala
          20             25             30

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ata gcg tct gac aag atc gct tcg acc ctc agg aga gga gga tgc tgt 144
 Ile Ala Ser Asp Lys Ile Ala Ser Thr Leu Arg Arg Gly Gly Cys Cys
 35 40 45

tct ttt cct gcc tgt aga aag tat cgt cca gaa atg tgt ggt gga cga 192
 Ser Phe Pro Ala Cys Arg Lys Tyr Arg Pro Glu Met Cys Gly Gly Arg
 50 55 60

cgc tgatgctcca ggaccctctg aaccacgacg t 226
 Arg
 65

<210> 93
 <211> 65
 <212> PRT
 <213> Conus sulcatus

<400> 93
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Phe Asn Ser Asp Arg Asp Pro Ala Leu Gly Gly Arg Asn Ala Ala Ala
 20 25 30

Ile Ala Ser Asp Lys Ile Ala Ser Thr Leu Arg Arg Gly Gly Cys Cys
 35 40 45

Ser Phe Pro Ala Cys Arg Lys Tyr Arg Pro Glu Met Cys Gly Gly Arg
 50 55 60

Arg
 65

<210> 94
 <211> 211
 <212> DNA
 <213> Conus sulcatus

<220>
 <221> CDS
 <222> (1)..(180)

<400> 94
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ttc act tca gat cat gaa tct gat cgc ggt gat gcc caa acc atc caa 96
 Phe Thr Ser Asp His Glu Ser Asp Arg Gly Asp Ala Gln Thr Ile Gln
 20 25 30

gaa gtg ttt gag atg ttc gct ctg gac agc gat gga tgc tgt tgg cat 144
 Glu Val Phe Glu Met Phe Ala Leu Asp Ser Asp Gly Cys Cys Trp His
 35 40 45

cct gct tgt ggc aga cac tat tgt ggt cga aga cgc tgatgctcca 190
 Pro Ala Cys Gly Arg His Tyr Cys Gly Arg Arg Arg
 50 55 60

ggaccctctg aaccacgacg t 211

<210> 95
 <211> 60
 <212> PRT
 <213> *Conus sulcatus*

<400> 95
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Thr Ser Asp His Glu Ser Asp Arg Gly Asp Ala Gln Thr Ile Gln
 20 25 30
 Glu Val Phe Glu Met Phe Ala Leu Asp Ser Asp Gly Cys Cys Trp His
 35 40 45
 Pro Ala Cys Gly Arg His Tyr Cys Gly Arg Arg Arg
 50 55 60

<210> 96
 <211> 202
 <212> DNA
 <213> *Conus sulcatus*

<220>
 <221> CDS
 <222> (1)..(195)

<400> 96
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc aat tca gat cgt gat cca gca tta ggt ggc agg aat gct gca gcc 96
 Phe Asn Ser Asp Arg Asp Pro Ala Leu Gly Gly Arg Asn Ala Ala Ala
 20 25 30
 ata gcg tct gac aag atc gct tcg acc ctc agg aga gga gga tgc tgt 144
 Ile Ala Ser Asp Lys Ile Ala Ser Thr Leu Arg Arg Gly Gly Cys Cys
 35 40 45
 tct ttt gct gcc tgt aga aag tat cgt cca gaa atg tgt ggt gga cga 192
 Ser Phe Ala Ala Cys Arg Lys Tyr Arg Pro Glu Met Cys Gly Gly Arg
 50 55 60
 cgc tgatgct 202
 Arg
 65

<210> 97
 <211> 65
 <212> PRT
 <213> *Conus sulcatus*

<400> 97
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Asn Ser Asp Arg Asp Pro Ala Leu Gly Gly Arg Asn Ala Ala Ala
 20 25 30
 Ile Ala Ser Asp Lys Ile Ala Ser Thr Leu Arg Arg Gly Gly Cys Cys
 35 40 45

Ser Phe Ala Ala Cys Arg Lys Tyr Arg Pro Glu Met Cys Gly Gly Arg
 50 55 60

Arg
 65

<210> 98
 <211> 220
 <212> DNA
 <213> Conus sulcatus

<220>
 <221> CDS
 <222> (1)..(189)

<400> 98
 atg ttc acc gtg ttt ctg ttg gtt ctc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Leu Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc aat tca gat cgt gca tta ggt ggc agg aat gct gca gcc aaa gcg 96
 Phe Asn Ser Asp Arg Ala Leu Gly Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 tct gac aag atc ctt tcg aac ctc agg aga gga gga tgc tgt ttt cat 144
 Ser Asp Lys Ile Leu Ser Asn Leu Arg Arg Gly Gly Cys Cys Phe His
 35 40 45
 cct gtc tgt tac atc aat ctt cta gaa atg tgt cgt caa cga ggc 189
 Pro Val Cys Tyr Ile Asn Leu Leu Glu Met Cys Arg Gln Arg Gly
 50 55 60
 tgatcgtcca ggaccctctg aaccacgacg t 220

<210> 99
 <211> 63
 <212> PRT
 <213> Conus sulcatus

<400> 99
 Met Phe Thr Val Phe Leu Leu Val Leu Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Asn Ser Asp Arg Ala Leu Gly Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 Ser Asp Lys Ile Leu Ser Asn Leu Arg Arg Gly Gly Cys Cys Phe His
 35 40 45
 Pro Val Cys Tyr Ile Asn Leu Leu Glu Met Cys Arg Gln Arg Gly
 50 55 60

<210> 100
 <211> 208
 <212> DNA
 <213> Conus consors

<220>
 <221> CDS
 <222> (1)..(177)

<400> 100

atg	ttc	acc	gtg	ttt	ctg	ttg	gtt	gtc	ttg	aca	acc	act	gtc	gtt	tcc	48
Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Thr	Thr	Thr	Val	Val	Ser	
1				5					10					15		

ttc	cct	tca	gat	agt	gca	tct	gat	gtc	agg	gat	gac	gaa	gcc	aaa	gac	96
Phe	Pro	Ser	Asp	Ser	Ala	Ser	Asp	Val	Arg	Asp	Asp	Glu	Ala	Lys	Asp	
			20					25					30			

gaa	agg	tct	gac	atg	tac	aaa	tcg	aaa	cgg	aat	gga	cgc	tgt	tgc	cat	144
Glu	Arg	Ser	Asp	Met	Tyr	Lys	Ser	Lys	Arg	Asn	Gly	Arg	Cys	Cys	His	
		35					40					45				

cct	gcc	tgt	ggc	aaa	cac	ttt	agt	tgt	gga	cgc	tgatgctcca	ggaccctctg	197
Pro	Ala	Cys	Gly	Lys	His	Phe	Ser	Cys	Gly	Arg			
	50					55							

aaccacgacg	t	208
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<210> 101

<211> 59

<212> PRT

<213> Conus consors

<400> 101

Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Thr	Thr	Thr	Val	Val	Ser
1				5					10					15	

Phe	Pro	Ser	Asp	Ser	Ala	Ser	Asp	Val	Arg	Asp	Asp	Glu	Ala	Lys	Asp
			20					25					30		

Glu	Arg	Ser	Asp	Met	Tyr	Lys	Ser	Lys	Arg	Asn	Gly	Arg	Cys	Cys	His
		35					40					45			

Pro	Ala	Cys	Gly	Lys	His	Phe	Ser	Cys	Gly	Arg
	50					55				

<210> 102

<211> 219

<212> DNA

<213> Conus stercusmuscarum

<220>

<221> CDS

<222> (1)..(189)

<400> 102

atg	ttc	acc	gtg	ttt	ctg	ttg	gtt	gtc	ttg	gca	acc	act	gtc	gtt	tcc	48
Met	Phe	Thr	Val	Phe	Leu	Leu	Val	Val	Leu	Ala	Thr	Thr	Val	Val	Ser	
1				5					10					15		

tcc	cct	tca	gat	cgt	gca	tct	gat	ggc	agg	aat	gcc	gca	gcc	aac	gag	96
Ser	Pro	Ser	Asp	Arg	Ala	Ser	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Asn	Glu	
			20					25					30			

aaa	gcg	tct	gac	gtg	atc	gcg	ctg	gcc	ctc	aag	gga	tgc	tgt	tcc	aac	144
Lys	Ala	Ser	Asp	Val	Ile	Ala	Leu	Ala	Leu	Lys	Gly	Cys	Cys	Ser	Asn	
		35					40					45				

cct	gtc	tgt	cac	ctg	gag	cat	tca	aac	atg	tgt	ggt	aga	aga	cgc		189
Pro	Val	Cys	His	Leu	Glu	His	Ser	Asn	Met	Cys	Gly	Arg	Arg	Arg		
	50					55					60					

tgatgctcca ggaccctctg aaccacgacg

219

<210> 103
 <211> 63
 <212> PRT
 <213> Conus stercusmuscarum

<400> 103
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Ser Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Glu
 20 25 30
 Lys Ala Ser Asp Val Ile Ala Leu Ala Leu Lys Gly Cys Cys Ser Asn
 35 40 45
 Pro Val Cys His Leu Glu His Ser Asn Met Cys Gly Arg Arg Arg
 50 55 60

<210> 104
 <211> 248
 <212> DNA
 <213> Conus betulinus

<220>
 <221> CDS
 <222> (1)..(180)

<400> 104
 atg ttc tcc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Ser Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 tcc act tca ggt ggt gca tct ggt ggc agg aag gct gca gcc aaa gcg 96
 Ser Thr Ser Gly Gly Ala Ser Gly Gly Arg Lys Ala Ala Ala Lys Ala
 20 25 30
 tct aac cgg atc gct ctg acc gtc agg agt gca aca tgc tgt aat tat 144
 Ser Asn Arg Ile Ala Leu Thr Val Arg Ser Ala Thr Cys Cys Asn Tyr
 35 40 45
 cct ccc tgt tac gag act tat cca gaa agt tgt ctg taacgtgaat 190
 Pro Pro Cys Tyr Glu Thr Tyr Pro Glu Ser Cys Leu
 50 55 60
 catccagagc ttgtgtggctg aagacactga tgctccagga ccctctgaac cacgacgt 248

<210> 105
 <211> 60
 <212> PRT
 <213> Conus betulinus

<400> 105
 Met Phe Ser Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Ser Thr Ser Gly Gly Ala Ser Gly Gly Arg Lys Ala Ala Ala Lys Ala
 20 25 30
 Ser Asn Arg Ile Ala Leu Thr Val Arg Ser Ala Thr Cys Cys Asn Tyr

35 40 45

Pro Pro Cys Tyr Glu Thr Tyr Pro Glu Ser Cys Leu
50 55 60

<210> 106
<211> 223
<212> DNA
<213> Conus betulinus

<220>
<221> CDS
<222> (1)..(183)

<400> 106
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtg gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc act tca ggt cgt gca ttt cgt ggc agg aat cgc gca gcc gac gac 96
Phe Thr Ser Gly Arg Ala Phe Arg Gly Arg Asn Arg Ala Ala Asp Asp
20 25 30

aaa agg tct gac ctg gcc gct ctg agc gtc agg gga gga tgc tgt tcc 144
Lys Arg Ser Asp Leu Ala Ala Leu Ser Val Arg Gly Gly Cys Cys Ser
35 40 45

cat cct gcc tgt gcg gtg aat cat cca gag ctt tgt ggc tgaagacgct 193
His Pro Ala Cys Ala Val Asn His Pro Glu Leu Cys Gly
50 55 60

gatgccccag gaccctctga accacgacgt 223

<210> 107
<211> 61
<212> PRT
<213> Conus betulinus

<400> 107
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Thr Ser Gly Arg Ala Phe Arg Gly Arg Asn Arg Ala Ala Asp Asp
20 25 30

Lys Arg Ser Asp Leu Ala Ala Leu Ser Val Arg Gly Gly Cys Cys Ser
35 40 45

His Pro Ala Cys Ala Val Asn His Pro Glu Leu Cys Gly
50 55 60

<210> 108
<211> 248
<212> DNA
<213> Conus betulinus

<220>
<221> CDS
<222> (1)..(180)

<400> 108

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc act tca ggt cgt gca tct ggt ggc agg aat gct gca gcc aaa gcg 96
Phe Thr Ser Gly Arg Ala Ser Gly Gly Arg Asn Ala Ala Ala Lys Ala
20 25 30

tct aac cgg atc gct atg gcc atc agc agt gga gca tgc tgt gca tat 144
Ser Asn Arg Ile Ala Met Ala Ile Ser Ser Gly Ala Cys Cys Ala Tyr
35 40 45

cct ccc tgt ttc gag gct tat cca gaa aga tgt ctg taacgtgaat 190
Pro Pro Cys Phe Glu Ala Tyr Pro Glu Arg Cys Leu
50 55 60

catccagacc tttgtggctg aagacgctga tgccccagga ccctctgaac cacgacgt 248

<210> 109
<211> 60
<212> PRT
<213> Conus betulinus

<400> 109
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Thr Ser Gly Arg Ala Ser Gly Gly Arg Asn Ala Ala Ala Lys Ala
20 25 30

Ser Asn Arg Ile Ala Met Ala Ile Ser Ser Gly Ala Cys Cys Ala Tyr
35 40 45

Pro Pro Cys Phe Glu Ala Tyr Pro Glu Arg Cys Leu
50 55 60

<210> 110
<211> 223
<212> DNA
<213> Conus betulinus

<220>
<221> CDS
<222> (1)..(192)

<400> 110
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc act tca gat cgt gca ttt cgt ggc agg aat tcc gca gcc aac gac 96
Phe Thr Ser Asp Arg Ala Phe Arg Gly Arg Asn Ser Ala Ala Asn Asp
20 25 30

aaa agg tct gac ctg gcc gct ctg agc gtc agg aga gga tgc tgc tcc 144
Lys Arg Ser Asp Leu Ala Ala Leu Ser Val Arg Arg Gly Cys Cys Ser
35 40 45

cat ccc gcc tgt agc gtg aat cat cca gag ctt tgt ggt aga aga cgc 192
His Pro Ala Cys Ser Val Asn His Pro Glu Leu Cys Gly Arg Arg Arg
50 55 60

tgatgccccca ggaccctctg aaccacgacg t

223

<210> 111
 <211> 64
 <212> PRT
 <213> Conus betulinus

<400> 111
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Thr Ser Asp Arg Ala Phe Arg Gly Arg Asn Ser Ala Ala Asn Asp
 20 25 30
 Lys Arg Ser Asp Leu Ala Ala Leu Ser Val Arg Arg Gly Cys Cys Ser
 35 40 45
 His Pro Ala Cys Ser Val Asn His Pro Glu Leu Cys Gly Arg Arg Arg
 50 55 60

<210> 112
 <211> 248
 <212> DNA
 <213> Conus betulinus

<220>
 <221> CDS
 <222> (1)..(180)

<400> 112
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc act tca ggt cgt gca tct ggt ggc agg aat gct gca gcc aaa gcg 96
 Phe Thr Ser Gly Arg Ala Ser Gly Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 tct aac cgg atc gct ctg atc gtc agg aat gca gaa tgc tgt tat tat 144
 Ser Asn Arg Ile Ala Leu Ile Val Arg Asn Ala Glu Cys Cys Tyr Tyr
 35 40 45
 cct ccc tgt tac gag gct tat cca gaa att tgt ctg taacgtgaat 190
 Pro Pro Cys Tyr Glu Ala Tyr Pro Glu Ile Cys Leu
 50 55 60
 catccagacc tttgtggctg aagaccctga tgctccagga ccctctgaac caccgacgt 248

<210> 113
 <211> 60
 <212> PRT
 <213> Conus betulinus

<400> 113
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Thr Ser Gly Arg Ala Ser Gly Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 Ser Asn Arg Ile Ala Leu Ile Val Arg Asn Ala Glu Cys Cys Tyr Tyr

35

40

45

Pro Pro Cys Tyr Glu Ala Tyr Pro Glu Ile Cys Leu
 50 55 60

<210> 114
 <211> 207
 <212> DNA
 <213> Conus pennaceus

<220>
 <221> CDS
 <222> (1)..(168)

<400> 114
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc att tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Ile Ser
 1 5 10 15
 ttc act tca gat cgt gca tct gat ggc ggg aat gcc gca gcg tct gac 96
 Phe Thr Ser Asp Arg Ala Ser Asp Gly Gly Asn Ala Ala Ala Ser Asp
 20 25 30
 ctg atc gct ctg acc atc aag gga tgc tgt tct cat cct ccc tgt gcc 144
 Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser His Pro Pro Cys Ala
 35 40 45
 atg aat aat cca gac tat tgt ggt tgacgacgct gatgctccag gaccctctga 198
 Met Asn Asn Pro Asp Tyr Cys Gly
 50 55
 accacgacg 207

<210> 115
 <211> 56
 <212> PRT
 <213> Conus pennaceus

<400> 115
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Ile Ser
 1 5 10 15
 Phe Thr Ser Asp Arg Ala Ser Asp Gly Gly Asn Ala Ala Ala Ser Asp
 20 25 30
 Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser His Pro Pro Cys Ala
 35 40 45
 Met Asn Asn Pro Asp Tyr Cys Gly
 50 55

<210> 116
 <211> 207
 <212> DNA
 <213> Conus pennaceus

<220>
 <221> CDS
 <222> (1)..(168)

<400> 116

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ttc act tca gat cgt gca tct gat ggc ggg aat gcc gca atg tct gac 96
 Phe Thr Ser Asp Arg Ala Ser Asp Gly Gly Asn Ala Ala Met Ser Asp
 20 25 30

ctg atc gct ctg acc atc aag gga tgc tgt tct cat cct ccc tgt ttc 144
 Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser His Pro Pro Cys Phe
 35 40 45

ctg aat aat cca gac tat tgt ggt tgacgacgct gatgctccag gaccctctga 198
 Leu Asn Asn Pro Asp Tyr Cys Gly
 50 55

accacgacg 207

<210> 117
 <211> 56
 <212> PRT
 <213> Conus pennaceus

<400> 117
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Phe Thr Ser Asp Arg Ala Ser Asp Gly Gly Asn Ala Ala Met Ser Asp
 20 25 30

Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser His Pro Pro Cys Phe
 35 40 45

Leu Asn Asn Pro Asp Tyr Cys Gly
 50 55

<210> 118
 <211> 210
 <212> DNA
 <213> Conus stercusmuscarum

<220>
 <221> CDS
 <222> (1)..(171)

<400> 118
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ttc cct tca gat cgt gaa tct gat ggc gcg aat gac gaa gcc cgc acc 96
 Phe Pro Ser Asp Arg Glu Ser Asp Gly Ala Asn Asp Glu Ala Arg Thr
 20 25 30

gac gag cct gag gag cac gga ccg gac agg aat gga tgc tgt agg aat 144
 Asp Glu Pro Glu Glu His Gly Pro Asp Arg Asn Gly Cys Cys Arg Asn
 35 40 45

cct gcc tgt gag agc cac aga tgt ggt tgacgacgct gatgctccag 191
 Pro Ala Cys Glu Ser His Arg Cys Gly
 50 55

gaccctctga accacgacg

210

<210> 119
 <211> 57
 <212> PRT
 <213> Conus stercusmuscarum

<400> 119
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Arg Glu Ser Asp Gly Ala Asn Asp Glu Ala Arg Thr
 20 25 30
 Asp Glu Pro Glu Glu His Gly Pro Asp Arg Asn Gly Cys Cys Arg Asn
 35 40 45
 Pro Ala Cys Glu Ser His Arg Cys Gly
 50 55

<210> 120
 <211> 210
 <212> DNA
 <213> Conus circumciscus

<220>
 <221> CDS
 <222> (1)..(180)

<400> 120
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc cct tca gat cgt gca tct gat ggc agg aat gcc gca gcc agc gac 96
 Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ser Asp
 20 25 30
 aga gcg tct gac gcg gcc cac cag gga tgc tgt tcc aac cct gtc tgt 144
 Arg Ala Ser Asp Ala Ala His Gln Gly Cys Cys Ser Asn Pro Val Cys
 35 40 45
 cac gtg gaa cat cca gaa ctt tgt cgt aga aga cgc tgatgctcca 190
 His Val Glu His Pro Glu Leu Cys Arg Arg Arg Arg
 50 55 60
 ggaccctctg aaccacgacg 210

<210> 121
 <211> 60
 <212> PRT
 <213> Conus circumciscus

<400> 121
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Ser Asp
 20 25 30
 Arg Ala Ser Asp Ala Ala His Gln Gly Cys Cys Ser Asn Pro Val Cys

35 40 45
 His Val Glu His Pro Glu Leu Cys Arg Arg Arg Arg
 50 55 60

<210> 122
 <211> 213
 <212> DNA
 <213> Conus circumciscus

<220>
 <221> CDS
 <222> (1)..(174)

<400> 122
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

 ttc cct tca aat cgt gaa tct gat ggc gcg aat gcc gaa gtc cgc acc 96
 Phe Pro Ser Asn Arg Glu Ser Asp Gly Ala Asn Ala Glu Val Arg Thr
 20 25 30

 gac gag cct gag gag cac gac gaa ctg ggc ggg aat gga tgc tgt ggg 144
 Asp Glu Pro Glu Glu His Asp Glu Leu Gly Gly Asn Gly Cys Cys Gly
 35 40 45

 aat cct gac tgt acg agc cac agt tgt gat tgacgacgct gatgctccag 194
 Asn Pro Asp Cys Thr Ser His Ser Cys Asp
 50 55

 gaccctctga accacgacg 213

<210> 123
 <211> 58
 <212> PRT
 <213> Conus circumciscus

<400> 123
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

 Phe Pro Ser Asn Arg Glu Ser Asp Gly Ala Asn Ala Glu Val Arg Thr
 20 25 30

 Asp Glu Pro Glu Glu His Asp Glu Leu Gly Gly Asn Gly Cys Cys Gly
 35 40 45

 Asn Pro Asp Cys Thr Ser His Ser Cys Asp
 50 55

<210> 124
 <211> 207
 <212> DNA
 <213> Conus episcopatus

<220>
 <221> CDS
 <222> (1)..(168)

<400> 124

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc act tca gat cgt gca tct gat agc agg aag gac gca gcg tct ggc 96
 Phe Thr Ser Asp Arg Ala Ser Asp Ser Arg Lys Asp Ala Ala Ser Gly
 20 25 30
 ctg atc gct ctg acc atc aag gga tgc tgt tct gat cct cgc tgt aac 144
 Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser Asp Pro Arg Cys Asn
 35 40 45
 atg aat aat cca gac tat tgt ggt tgacgacgct gatgctccag gaccctctga 198
 Met Asn Asn Pro Asp Tyr Cys Gly
 50 55
 accacgacg 207

<210> 125
 <211> 56
 <212> PRT
 <213> *Conus episcopatus*

<400> 125
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Thr Ser Asp Arg Ala Ser Asp Ser Arg Lys Asp Ala Ala Ser Gly
 20 25 30
 Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser Asp Pro Arg Cys Asn
 35 40 45
 Met Asn Asn Pro Asp Tyr Cys Gly
 50 55

<210> 126
 <211> 213
 <212> DNA
 <213> *Conus sponsalis*

<220>
 <221> CDS
 <222> (1)..(174)

<400> 126
 atg tcc acc gtg ttt ctg ttg gtt gtc ctc gca acc acc gtc gtt tcc 48
 Met Ser Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc act gta gat cgt gca tct gat ggc agg gat gtc gca atc gac gac 96
 Phe Thr Val Asp Arg Ala Ser Asp Gly Arg Asp Val Ala Ile Asp Asp
 20 25 30
 aga ttg gtg tct ctc cct cag atc gcc cat gct gac tgt tgt tcc gat 144
 Arg Leu Val Ser Leu Pro Gln Ile Ala His Ala Asp Cys Cys Ser Asp
 35 40 45
 cct gcc tgc aag cag acg ccc ggt tgt cgt taaagacgct gctgctccag 194
 Pro Ala Cys Lys Gln Thr Pro Gly Cys Arg
 50 55

gaccctctga accacgacg

213

<210> 127

<211> 58

<212> PRT

<213> Conus sponsalis

<400> 127

Met Ser Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Phe Thr Val Asp Arg Ala Ser Asp Gly Arg Asp Val Ala Ile Asp Asp
 20 25 30

Arg Leu Val Ser Leu Pro Gln Ile Ala His Ala Asp Cys Cys Ser Asp
 35 40 45

Pro Ala Cys Lys Gln Thr Pro Gly Cys Arg
 50 55

<210> 128

<211> 221

<212> DNA

<213> Conus sponsalis

<220>

<221> CDS

<222> (1)..(168)

<400> 128

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gct tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Ala Ser
 1 5 10 15

ttc att atc gat gat cca tct gat ggc agg aat att gca gtc gac gac 96
 Phe Ile Ile Asp Asp Pro Ser Asp Gly Arg Asn Ile Ala Val Asp Asp
 20 25 30

aga ggg ctt ttc tct acg ctc ttc cat gct gat tgc tgt gaa aat cct 144
 Arg Gly Leu Phe Ser Thr Leu Phe His Ala Asp Cys Cys Glu Asn Pro
 35 40 45

gcc tgt aga cac acg cag ggt tgt tgatctttgt tcttcaaaga cactgctggc 198
 Ala Cys Arg His Thr Gln Gly Cys
 50 55

ccaggaccct ctgaaccacg acg

221

<210> 129

<211> 56

<212> PRT

<213> Conus sponsalis

<400> 129

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Ala Ser
 1 5 10 15

Phe Ile Ile Asp Asp Pro Ser Asp Gly Arg Asn Ile Ala Val Asp Asp
 20 25 30

Arg Gly Leu Phe Ser Thr Leu Phe His Ala Asp Cys Cys Glu Asn Pro

35 40 45

Ala Cys Arg His Thr Gln Gly Cys
50 55

<210> 130
<211> 220
<212> DNA
<213> Conus dalli

<220>
<221> CDS
<222> (1)..(180)

<400> 130
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc act tca gat cgt gca ttt cgt ggc agg aat gcc gca gcc aaa gag 96
Phe Thr Ser Asp Arg Ala Phe Arg Gly Arg Asn Ala Ala Ala Lys Glu
20 25 30

tct ggc ctg gtc ggt ctg acc gac aag acg cga gga tgc tgt tct cat 144
Ser Gly Leu Val Gly Leu Thr Asp Lys Thr Arg Gly Cys Cys Ser His
35 40 45

cct gcc tgt aac gta gat cat cca gaa att tgt ggt tgaagacgct 190
Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Gly
50 55 60

gatgctccag gaccctctga accacgacgt 220

<210> 131
<211> 60
<212> PRT
<213> Conus dalli

<400> 131
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Thr Ser Asp Arg Ala Phe Arg Gly Arg Asn Ala Ala Ala Lys Glu
20 25 30

Ser Gly Leu Val Gly Leu Thr Asp Lys Thr Arg Gly Cys Cys Ser His
35 40 45

Pro Ala Cys Asn Val Asp His Pro Glu Ile Cys Gly
50 55 60

<210> 132
<211> 208
<212> DNA
<213> Conus dalli

<220>
<221> CDS
<222> (1)..(177)

<400> 132

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc act tca gat ggt gca tct gat gac agg aaa gcc gct gcg tct gac 96
Phe Thr Ser Asp Gly Ala Ser Asp Asp Arg Lys Ala Ala Ala Ser Asp
20 25 30

ctg atc act ctg acc atc aag gga tgc tgt tct cgt cct ccc tgt atc 144
Leu Ile Thr Leu Thr Ile Lys Gly Cys Cys Ser Arg Pro Pro Cys Ile
35 40 45

gcg aat aat cca gac ttg tgt ggt cga cga cgc tgatgctcca ggaccctctg 197
Ala Asn Asn Pro Asp Leu Cys Gly Arg Arg Arg
50 55

aaccacgacg t 208

<210> 133
<211> 59
<212> PRT
<213> Conus dalli

<400> 133
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Thr Ser Asp Gly Ala Ser Asp Asp Arg Lys Ala Ala Ala Ser Asp
20 25 30

Leu Ile Thr Leu Thr Ile Lys Gly Cys Cys Ser Arg Pro Pro Cys Ile
35 40 45

Ala Asn Asn Pro Asp Leu Cys Gly Arg Arg Arg
50 55

<210> 134
<211> 223
<212> DNA
<213> Conus dalli

<220>
<221> CDS
<222> (1)..(192)

<400> 134
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

tcc act tca ggt cgt cgt gca ttt cat ggc agg aat gcc gca gcc aaa 96
Ser Thr Ser Gly Arg Arg Ala Phe His Gly Arg Asn Ala Ala Ala Lys
20 25 30

gcg tct gga ctg gtc ggt ctg act gac agg aga cca caa tgc tgt agt 144
Ala Ser Gly Leu Val Gly Leu Thr Asp Arg Arg Pro Gln Cys Cys Ser
35 40 45

gat cct cgc tgt aac gta ggt cat cca gaa ctt tgt ggt gga aga cgc 192
Asp Pro Arg Cys Asn Val Gly His Pro Glu Leu Cys Gly Gly Arg Arg
50 55 60

tgatgctcca ggaccctctg aaccacaacg t

223

<210> 135
 <211> 64
 <212> PRT
 <213> Conus dalli

<400> 135
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Ser Thr Ser Gly Arg Arg Ala Phe His Gly Arg Asn Ala Ala Ala Lys
 20 25 30
 Ala Ser Gly Leu Val Gly Leu Thr Asp Arg Arg Pro Gln Cys Cys Ser
 35 40 45
 Asp Pro Arg Cys Asn Val Gly His Pro Glu Leu Cys Gly Gly Arg Arg
 50 55 60

<210> 136
 <211> 220
 <212> DNA
 <213> Conus dalli

<220>
 <221> CDS
 <222> (1)..(189)

<400> 136
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 tcc act tca ggt cgt gca ttt cat ggc agg aat gcc gca gcc aaa gcg 96
 Ser Thr Ser Gly Arg Ala Phe His Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 tct ggc ctg gtc ggt ctg acc gac aag agg caa gta tgc tgt agt gat 144
 Ser Gly Leu Val Gly Leu Thr Asp Lys Arg Gln Val Cys Cys Ser Asp
 35 40 45
 cct cgc tgt aac gta ggt cat cca gaa att tgt ggt gga aga cgc 189
 Pro Arg Cys Asn Val Gly His Pro Glu Ile Cys Gly Gly Arg Arg
 50 55 60
 tgatgctcca ggaccctctg aaccacgacg t 220

<210> 137
 <211> 63
 <212> PRT
 <213> Conus dalli

<400> 137
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Ser Thr Ser Gly Arg Ala Phe His Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30
 Ser Gly Leu Val Gly Leu Thr Asp Lys Arg Gln Val Cys Cys Ser Asp

35

40

45

Pro Arg Cys Asn Val Gly His Pro Glu Ile Cys Gly Gly Arg Arg
 50 55 60

<210> 138
 <211> 208
 <212> DNA
 <213> *Conus achatinus*

<220>
 <221> CDS
 <222> (1)..(180)

<400> 138
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg aca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Thr Thr Thr Val Val Ser
 1 5 10 15
 ttc cct tca gat agt gca tct ggt ggc agg gat gac gag gcc aaa gac 96
 Phe Pro Ser Asp Ser Ala Ser Gly Gly Arg Asp Asp Glu Ala Lys Asp
 20 25 30
 gaa agg tct gac atg tac gaa ttg aaa cgg aat gga cgc tgt tgc cat 144
 Glu Arg Ser Asp Met Tyr Glu Leu Lys Arg Asn Gly Arg Cys Cys His
 35 40 45
 cct gcc tgt ggt ggc aaa tac gtt aaa tgt gga cgc tgatgctcca 190
 Pro Ala Cys Gly Gly Lys Tyr Val Lys Cys Gly Arg
 50 55 60
 ggaccctctc gaaccacg 208

<210> 139
 <211> 60
 <212> PRT
 <213> *Conus achatinus*

<400> 139
 Met Phe Thr Val Phe Leu Leu Val Val Leu Thr Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Ser Ala Ser Gly Gly Arg Asp Asp Glu Ala Lys Asp
 20 25 30
 Glu Arg Ser Asp Met Tyr Glu Leu Lys Arg Asn Gly Arg Cys Cys His
 35 40 45
 Pro Ala Cys Gly Gly Lys Tyr Val Lys Cys Gly Arg
 50 55 60

<210> 140
 <211> 211
 <212> DNA
 <213> *Conus bullatus*

<220>
 <221> CDS
 <222> (1)..(174)

<400> 140

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc tct aca gat gat gaa tct gat ggc tcg aat gaa gaa ccc agc gcc 96
Phe Ser Thr Asp Asp Glu Ser Asp Gly Ser Asn Glu Glu Pro Ser Ala
20 25 30

gac cag act gcc agg tcc tca atg aac agg gcg cct gga tgc tgt aac 144
Asp Gln Thr Ala Arg Ser Ser Met Asn Arg Ala Pro Gly Cys Cys Asn
35 40 45

aat cct gcc tgt gtg aag cac aga tgt gga tgacgctgat gctccaggac 194
Asn Pro Ala Cys Val Lys His Arg Cys Gly
50 55

cctctgaacc acgacgt 211

<210> 141
<211> 58
<212> PRT
<213> Conus bullatus

<400> 141
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Ser Thr Asp Asp Glu Ser Asp Gly Ser Asn Glu Glu Pro Ser Ala
20 25 30

Asp Gln Thr Ala Arg Ser Ser Met Asn Arg Ala Pro Gly Cys Cys Asn
35 40 45

Asn Pro Ala Cys Val Lys His Arg Cys Gly
50 55

<210> 142
<211> 214
<212> DNA
<213> Conus bullatus

<220>
<221> CDS
<222> (1)..(177)

<400> 142
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc tct aca gat gat gaa tct gat ggc tcg aat gaa gaa ccc agc gcc 96
Phe Ser Thr Asp Asp Glu Ser Asp Gly Ser Asn Glu Glu Pro Ser Ala
20 25 30

gac cag gct gcc agg tcc gca atg aac agg ccg cct gga tgc tgt aac 144
Asp Gln Ala Ala Arg Ser Ala Met Asn Arg Pro Pro Gly Cys Cys Asn
35 40 45

aat cct gcc tgt gtg aag cac aga tgt ggt gga tgacgctgat gctccaggac 197
Asn Pro Ala Cys Val Lys His Arg Cys Gly Gly
50 55

cctctgaacc acgacgt

214

<210> 143
 <211> 59
 <212> PRT
 <213> Conus bullatus

<400> 143
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Ser Thr Asp Asp Glu Ser Asp Gly Ser Asn Glu Glu Pro Ser Ala
 20 25 30
 Asp Gln Ala Ala Arg Ser Ala Met Asn Arg Pro Pro Gly Cys Cys Asn
 35 40 45
 Asn Pro Ala Cys Val Lys His Arg Cys Gly Gly
 50 55

<210> 144
 <211> 208
 <212> DNA
 <213> Conus bullatus

<220>
 <221> CDS
 <222> (1)..(177)

<400> 144
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 ttc cct tca gat cgt gac tct gat ggc gcg gat gcc gaa gcc agt gac 96
 Phe Pro Ser Asp Arg Asp Ser Asp Gly Ala Asp Ala Glu Ala Ser Asp
 20 25 30
 gag cct gtt gag ttc gaa agg gac gag aat gga tgc tgt tgg aat cct 144
 Glu Pro Val Glu Phe Glu Arg Asp Gly Asn Gly Cys Cys Trp Asn Pro
 35 40 45
 tcc tgt ccg agg ccc aga tgt aca gga cga cgc taatgctcca ggacctctg 197
 Ser Cys Pro Arg Pro Arg Cys Thr Gly Arg Arg
 50 55

aaccacgacg t

208

<210> 145
 <211> 59
 <212> PRT
 <213> Conus bullatus

<400> 145
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Arg Asp Ser Asp Gly Ala Asp Ala Glu Ala Ser Asp
 20 25 30
 Glu Pro Val Glu Phe Glu Arg Asp Gly Asn Gly Cys Cys Trp Asn Pro

35

40

45

Ser Cys Pro Arg Pro Arg Cys Thr Gly Arg Arg
 50 55

<210> 146
 <211> 211
 <212> DNA
 <213> Conus bullatus

<220>
 <221> CDS
 <222> (1)..(180)

<400> 146
 atg ttc acc gtg ttt ctg ttg gtt gtc ttg aca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Thr Thr Thr Val Val Ser
 1 5 10 15
 ttc cct tca gat cgt gca tct gat ggc agg aat gcc gca gcc aac gac 96
 Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp
 20 25 30
 aaa gcg tct gac gtg gtc acg ctg gtc ctc aag gga tgc tgt tcc acc 144
 Lys Ala Ser Asp Val Val Thr Leu Val Leu Lys Gly Cys Cys Ser Thr
 35 40 45
 cct ccc tgt gct gtg ctg tat tgt ggt aga aga cgc tgatgctcca 190
 Pro Pro Cys Ala Val Leu Tyr Cys Gly Arg Arg Arg
 50 55 60
 ggaccctctg aaccacgacg t 211

<210> 147
 <211> 60
 <212> PRT
 <213> Conus bullatus

<400> 147
 Met Phe Thr Val Phe Leu Leu Val Val Leu Thr Thr Thr Val Val Ser
 1 5 10 15
 Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp
 20 25 30
 Lys Ala Ser Asp Val Val Thr Leu Val Leu Lys Gly Cys Cys Ser Thr
 35 40 45
 Pro Pro Cys Ala Val Leu Tyr Cys Gly Arg Arg Arg
 50 55 60

<210> 148
 <211> 212
 <212> DNA
 <213> Conus distans

<220>
 <221> CDS
 <222> (1)..(171)

<400> 148

atg ttc acc gtg ttt ctg ttg gtt gtc ttc gca tcc tct gtc acc tta 48
Met Phe Thr Val Phe Leu Leu Val Val Phe Ala Ser Ser Val Thr Leu
1 5 10 15

gat cgt gca tct tat ggc agg tat gcc tca ccc gtc gac aga gcg tct 96
Asp Arg Ala Ser Tyr Gly Arg Tyr Ala Ser Pro Val Asp Arg Ala Ser
20 25 30

gcc ctg atc gct cag gcc atc ctt cga gat tgc tgc tcc aat cct cct 144
Ala Leu Ile Ala Gln Ala Ile Leu Arg Asp Cys Cys Ser Asn Pro Pro
35 40 45

tgt gcc cat aat aat cca gac tgt cgt taaagacgct gcttgctcca 191
Cys Ala His Asn Asn Pro Asp Cys Arg
50 55

ggaccctctg aaccacgacg t 212

<210> 149
<211> 57
<212> PRT
<213> Conus distans

<400> 149
Met Phe Thr Val Phe Leu Leu Val Val Phe Ala Ser Ser Val Thr Leu
1 5 10 15

Asp Arg Ala Ser Tyr Gly Arg Tyr Ala Ser Pro Val Asp Arg Ala Ser
20 25 30

Ala Leu Ile Ala Gln Ala Ile Leu Arg Asp Cys Cys Ser Asn Pro Pro
35 40 45

Cys Ala His Asn Asn Pro Asp Cys Arg
50 55

<210> 150
<211> 63
<212> DNA
<213> Conus textile

<220>
<221> CDS
<222> (1)..(60)

<400> 150
gga tgc tgt tct aat cct ccc tgt atc gcg aag aat cca cac atg tgt 48
Gly Cys Cys Ser Asn Pro Pro Cys Ile Ala Lys Asn Pro His Met Cys
1 5 10 15

ggt gga aga cgc tga 63
Gly Gly Arg Arg
20

<210> 151
<211> 20
<212> PRT
<213> Conus textile

<400> 151
Gly Cys Cys Ser Asn Pro Pro Cys Ile Ala Lys Asn Pro His Met Cys

1 5 10 15

Gly Gly Arg Arg
20

<210> 152
<211> 220
<212> DNA
<213> Conus consors

<220>
<221> CDS
<222> (1)..(189)

<400> 152
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

ttc cct tca gat cgt gca tct gat ggc agg aat gcc gca gcc aac gac 96
Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp
20 25 30

aaa gcg tct gac gtg atc acg ctg gcc ctc aag gga tgc tgt tcc aac 144
Lys Ala Ser Asp Val Ile Thr Leu Ala Leu Lys Gly Cys Cys Ser Asn
35 40 45

cct gtc tgt cac ttg gag cat tca aac ctt tgt ggt aga aga cgc 189
Pro Val Cys His Leu Glu His Ser Asn Leu Cys Gly Arg Arg Arg
50 55 60

tgatgctcca ggaccctctg aaccacgacg t 220

<210> 153
<211> 63
<212> PRT
<213> Conus consors

<400> 153
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15

Phe Pro Ser Asp Arg Ala Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp
20 25 30

Lys Ala Ser Asp Val Ile Thr Leu Ala Leu Lys Gly Cys Cys Ser Asn
35 40 45

Pro Val Cys His Leu Glu His Ser Asn Leu Cys Gly Arg Arg Arg
50 55 60

<210> 154
<211> 15
<212> PRT
<213> Conus musicus

<220>
<221> SITE
<222> (4)..(12)
<223> Xaa at residues 4, 11 and 12 is Tyr, nor-Tyr,
mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,

O-phospho-Tyr or nitro-Tyr. Xaa at residue 6 is Pro or hydroxy-Pro.

```
<220>
<221> SITE
<222> (9)..(15)
<223> Xaa at residues 9, 10 and 15 is Lys, N-methyl-Lys,
N,N-dimethyl-Lys or N,N,N-trimethyl-Lys; Xaa at
residue 14 is Trp (D or L) or halo-Trp.
```

<400> 154
Gly Cys Cys Xaa Asn Xaa Val Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10 15

<210> 155
<211> 16
<212> PRT
<213> Conus purpurascens

```
<220>
<221> SITE
<222> (1)..(3)
<223> Xaa at residue 1 is Gln or pyro-Glu; Xaa at
      residue 2 is Glu or gamma-carboxy-Glu; Xaa at
      residues 3 and 9 is Pro or hydroxy-Pro.
```

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<220>
<221> SITE
<222> (13)
<223> Xaa at residue 13 is Lys, N-methyl-Lys,
N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.
```

<400> 155
Xaa Xaa Xaa Gly Cys Cys Arg His Xaa Ala Cys Gly Xaa Asn Arg Cys
1 5 10 15

```
<210> 156
<211> 13
<212> PRT
<213> Conus musicus
```

```
<220>
<221> SITE
<222> (5)..(11)
<223> Xaa at residues 5 and 11 is Pro or hydroxy-Pro.
```

<400> 156
Cys Cys Ala Asp Xaa Asp Cys Arg Phe Arg Xaa Gly Cys
1 5 10

```
<210> 157
<211> 17
<212> PRT
<213> Conus musicus
```

```
<220>
<221> SITE
<222> (4)..(13)
<223> Xaa at residues 4 and 13 is Tyr, nor-Tyr,
      mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
      O-phospho-Tyr or nitro-Tyr; Xaa at residues 6 and
```

10 is Pro or hydroxy-Pro.

<220>
 <221> SITE
 <222> (9)..(17)
 <223> Xaa at residues 9 and 16 is Trp (D or L) or
 halo-Trp; Xaa at residues 11 and 17 is Lys,
 N-methyl-Lys, N,N-dimethyl-Lys or
 N,N,N-trimethyl-Lys.
 <400> 157
 Gly Cys Cys Xaa Asn Xaa Ser Cys Xaa Xaa Xaa Thr Xaa Cys Ser Xaa
 1 5 10 15

Xaa

<210> 158
 <211> 13
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (5)..(8)
 <223> Xaa at residue 5 is Pro or hydroxy-Pro; Xaa at
 residue 8 is Lys, N-methyl-Lys, N,N-dimethyl-Lys
 or N,N,N-trimethyl-Lys.

<220>
 <221> SITE
 <222> (9)..(11)
 <223> Xaa at residue 9 is Glu or gamma-carboxy-Glu; Xaa
 at residue 11 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 158
 Cys Cys Ser Asn Xaa Thr Cys Xaa Xaa Thr Xaa Gly Cys
 1 5 10

<210> 159
 <211> 13
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (5)..(11)
 <223> Xaa at residues 5 and 11 is Pro or hydroxy-Pro;
 Xaa at residue 8 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 159
 Cys Cys Ala Asn Xaa Ile Cys Xaa Asn Thr Xaa Gly Cys
 1 5 10

<210> 160
 <211> 13
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (5)..(8)
 <223> Xaa at residue 5 is Pro or hydroxy-Pro; Xaa at residue 8 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<220>
 <221> SITE
 <222> (9)..(11)
 <223> Xaa at residue 9 is Glu or gamma-carboxy-Glu; Xaa at residue 11 is Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<400> 160
 Cys Cys Asn Asn Xaa Thr Cys Xaa Xaa Thr Xaa Gly Cys
 1 5 10

<210> 161
 <211> 13
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (5)..(8)
 <223> Xaa at residue 5 is Pro or hydroxy-Pro; Xaa at residue 8 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<220>
 <221> SITE
 <222> (9)..(11)
 <223> Xaa at residue 9 is Glu or gamma-carboxy-Glu; Xaa at residue 11 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<400> 161
 Cys Cys Ser Asn Xaa Val Cys Xaa Xaa Thr Xaa Gly Cys
 1 5 10

<210> 162
 <211> 17
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residue 6 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr; Xaa at residues 7, 8 and 14 is Pro or hydroxy-Pro.

<220>
 <221> SITE
 <222> (15)
 <223> Xaa at residue 15 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 162

Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile Ala Ser Asn Xaa Xaa Cys
 1 5 10 15

Gly

<210> 163
 <211> 15
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 163
 Gly Cys Cys Ser His Xaa Val Cys Ser Ala Met Ser Xaa Ile Cys
 1 5 10 15

<210> 164
 <211> 15
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (4)..(12)
 <223> Xaa at residues 4 and 12 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys; Xaa at
 residue 6 is Pro or hydroxy-Pro.

<220>
 <221> SITE
 <222> (7)..(14)
 <223> Xaa at residues 7 and 14 is Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 164
 Gly Cys Cys Xaa Asn Xaa Xaa Cys Gly Ala Ser Xaa Thr Xaa Cys
 1 5 10 15

<210> 165
 <211> 15
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (5)..(13)
 <223> Xaa at residue 5 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr; Xaa at residues 6, 7 and 13 is Pro or
 hydroxy-Pro.

<400> 165
 Gly Cys Cys Ser Xaa Xaa Xaa Cys Phe Ala Thr Asn Xaa Asp Cys
 1 5 10 15

<210> 166
 <211> 17
 <212> PRT
 <213> Conus radiatus

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residue 6 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr; Xaa at residues 7, 8 and 14 is Pro or hydroxy-Pro.

<400> 166
 Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile Ala Asn Asn Xaa Leu Cys
 1 5 10 15

Ala

<210> 167
 <211> 17
 <212> PRT
 <213> Conus radiatus

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residue 6 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr; Xaa at residues 7, 8 and 14 is Pro or hydroxy-Pro.

<400> 167
 Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile Ala Asn Asn Xaa Phe Cys
 1 5 10 15

Ala

<210> 168
 <211> 16
 <212> PRT
 <213> Conus virgo

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro.

<400> 168
 Asp Cys Cys Ser Asn Xaa Xaa Cys Ser Gln Asn Asn Xaa Asp Cys Met
 1 5 10 15

<210> 169
 <211> 16
 <212> PRT
 <213> Conus virgo

<220>
 <221> SITE

<222> (6)..(13)

<223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro.

<400> 169

Asp Cys Cys Ser Asn Xaa Xaa Cys Ala His Asn Asn Xaa Asp Cys Arg
 1 5 10 15

<210> 170

<211> 20

<212> PRT

<213> Conus achatinus

<220>

<221> SITE

<222> (1)..(14)

<223> Xaa at residues 1, 11 and 14 is Glu or
 gamma-carboxy-Glu; Xaa at residue 6 is Pro or
 hydroxy-Pro.

<400> 170

Xaa Cys Cys Thr Asn Xaa Val Cys His Ala Xaa His Gln Xaa Leu Cys
 1 5 10 15

Ala Arg Arg Arg
 20

<210> 171

<211> 16

<212> PRT

<213> Conus achatinus

<220>

<221> SITE

<222> (6)..(10)

<223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residue 11 is Glu or gamma-carboxy-Glu.

<400> 171

Gly Cys Cys Ser Asn Xaa Val Cys His Leu Xaa His Ser Asn Leu Cys
 1 5 10 15

<210> 172

<211> 20

<212> PRT

<213> Conus achatinus

<220>

<221> SITE

<222> (1)..(14)

<223> Xaa at residues 1, 11 and 14 is Glu or
 gamma-carboxy-Glu; Xaa at residue 6 is Pro or
 hydroxy-Pro.

<400> 172

Xaa Cys Cys Thr Asn Xaa Val Cys His Val Xaa His Gln Xaa Leu Cys
 1 5 10 15

Ala Arg Arg Arg
 20

<210> 173
 <211> 17
 <212> PRT
 <213> *Conus ammiralis*

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residue 1 is Gln or pyro-Glu; Xaa at residues 2 and 15 is Glu or gamma-carboxy-Glu; Xaa at residue 6 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>
 <221> SITE
 <222> (6)..(14)
 <223> nitro-Tyr; Xaa at residues 7 and 14 is Pro or hydroxy-Pro.

<400> 173
 Xaa Xaa Cys Cys Ser Xaa Xaa Ala Cys Asn Leu Asp His Xaa Xaa Leu
 1 5 10 15

Cys

<210> 174
 <211> 18
 <212> PRT
 <213> *Conus ammiralis*

<220>
 <221> SITE
 <222> (1)..(15)
 <223> Xaa at residues 1, 7 and 14 is Pro or hydroxy-Pro; Xaa at residues 2 and 15 is Glu or gamma-carboxy-Glu.

<400> 174
 Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Asn Ser Thr His Xaa Xaa Leu
 1 5 10 15

Cys Gly

<210> 175
 <211> 21
 <212> PRT
 <213> *Conus arenatus*

<220>
 <221> SITE
 <222> (7)..(12)
 <223> Xaa at residues 7 and 8 is Pro or hydroxy-Pro; Xaa at residue 10 is Trp (D or L) or halo-Trp; Xaa at residues 11 and 12 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<220>
 <221> SITE
 <222> (13)..(19)

<223> Xaa at residue 13 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr ; Xaa at residue 19 is Glu or gamma-carboxy-Glu.

<400> 175

Leu Asn Cys Cys Met Ile Xaa Xaa Cys Xaa Xaa Xaa Xaa Gly Asp Arg
1 5 10 15

Cys Ser Xaa Val Arg
20

<210> 176

<211> 22

<212> PRT

<213> Conus arenatus

<220>

<221> SITE

<222> (9)..(20)

<223> Xaa at residue 9 is Pro or hydroxy-Pro; Xaa at residues 12 and 20 is Glu or gamma-carboxy-Glu; Xaa at residue 14 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>

<221> SITE

<222> (9)..(20)

<223> nitro-Tyr.

<400> 176

Ala Phe Gly Cys Cys Asp Leu Ile Xaa Cys Leu Xaa Arg Xaa Gly Asn
1 5 10 15

Arg Cys Asn Xaa Val His
20

<210> 177

<211> 21

<212> PRT

<213> Conus arenatus

<220>

<221> SITE

<222> (8)..(16)

<223> Xaa at residue 8 is Pro or hydroxy-Pro; Xaa at residue 10 is Trp (D or L) or halo-Trp; Xaa at residues 12 and 16 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<220>

<221> SITE

<222> (11)..(19)

<223> Xaa at residues 11 and 19 is Glu or gamma-carboxy-Glu; Xaa at residue 13 is Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<400> 177

Leu Gly Cys Cys Asn Val Thr Xaa Cys Xaa Xaa Xaa Xaa Gly Asp Xaa
1 5 10 15

Cys Asn Xaa Val Arg
20

<210> 178
<211> 20
<212> PRT
<213> Conus arenatus

<220>
<221> SITE
<222> (2)..(14)
<223> Xaa at residue 2 is Glu or gamma-carboxy-Glu; Xaa
at residues 7 and 14 is Pro or hydroxy-Pro.

<400> 178
Asp Xaa Cys Cys Ser Asn Xaa Ala Cys Arg Val Asn Asn Xaa His Val
1 5 10 15

Cys Arg Arg Arg
20

<210> 179
<211> 21
<212> PRT
<213> Conus arenatus

<220>
<221> SITE
<222> (7)..(13)
<223> Xaa at residue 7 is Pro or hydroxy-Pro; Xaa at
residue 10 is Trp (D or L) or halo-Trp; Xaa at
residue 12 is Glu or gamma-carboxy-Glu; Xaa at
residue 13 is Tyr, nor-Tyr, mono-halo-Tyr,

<220>
<221> SITE
<222> (13)..(19)
<223> di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
nitro-Tyr; Xaa at residues 14 and 19 is Lys,
N-methyl-Lys, N,N-dimethyl-Lys or
N,N,N-trimethyl-Lys.

<400> 179
Leu Asn Cys Cys Ser Ile Xaa Gly Cys Xaa Asn Xaa Xaa Xaa Asp Arg
1 5 10 15

Cys Ser Xaa Val Arg
20

<210> 180
<211> 18
<212> PRT
<213> Conus aurisiacus

<220>
<221> SITE
<222> (7)..(14)
<223> Xaa at residues 7 and 14 is Pro or hydroxy-Pro;
Xaa at residue 10 is Tyr, mono-halo-Tyr,
di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
nitro-Tyr.

<400> 180
 Gly Gly Cys Cys Ser His Xaa Val Cys Xaa Phe Asn Asn Xaa Gln Met
 1 5 10 15

Cys Arg

<210> 181
 <211> 18
 <212> PRT
 <213> Conus aurisiacus

<220>
 <221> SITE
 <222> (7)..(14)
 <223> Xaa at residues 7 and 14 is Pro or hydroxy-Pro.

<400> 181
 Gly Gly Cys Cys Ser His Xaa Val Cys Asn Leu Asn Asn Xaa Gln Met
 1 5 10 15

Cys Arg

<210> 182
 <211> 17
 <212> PRT
 <213> Conus bandanus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residues 6 and 7 is Pro or hydroxy-Pro; Xaa
 at residues 9 and 15 is Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 182
 Gly Cys Cys Ser His Xaa Xaa Cys Xaa Ala Asn Asn Gln Ala Xaa Cys
 1 5 10 15

Asn

<210> 183
 <211> 17
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (7)..(15)
 <223> Xaa at residues 7 and 14 is Pro and hydroxy-Pro;
 Xaa at residue 15 is Glu or gamma-carboxy-Glu.

<400> 183
 Gly Gly Cys Cys Ser His Xaa Ala Cys Ser Val Thr His Xaa Xaa Leu
 1 5 10 15

Cys

<210> 184
 <211> 18
 <212> PRT
 <213> Conus betulinus

<220>
 <221> SITE
 <222> (6)..(12)
 <223> Xaa at residue 6 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr; Xaa at residue 7 is Pro and hydroxy-Pro; Xaa at residue 12 is Glu or

<220>
 <221> SITE
 <222> (6)..(12)
 <223> gamma-carboxy-Glu.

<400> 184
 Gly Gly Cys Cys Ser Xaa Xaa Ala Cys Ser Val Xaa His Gln Asp Leu
 1 5 10 15

Cys Asp

<210> 185
 <211> 25
 <212> PRT
 <213> Conus characteristicus

<220>
 <221> SITE
 <222> (8)..(22)
 <223> Xaa at residues 8 and 22 is Pro or hydroxy-Pro; Xaa at residue 10 is Trp (D or L) or halo-Trp; Xaa at residue 13 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>
 <221> SITE
 <222> (8)..(22)
 <223> nitro-Tyr; Xaa at residues 15, 16 and 19 is Glu or gamma-carboxy-Glu.

<400> 185
 Val Ser Cys Cys Val Val Arg Xaa Cys Xaa Ile Arg Xaa Gln Xaa Xaa
 1 5 10 15

Cys Leu Xaa Ala Asp Xaa Arg Thr Leu
 20 25

<210> 186
 <211> 21
 <212> PRT
 <213> Conus characteristicus

<220>
 <221> SITE
 <222> (1)..(19)
 <223> Xaa at residue 1 is Gln or pyro-Glu; Xaa at

residue 7 is Pro or hydroxy-Pro; Xaa at residue 10 is Trp (D or L) or halo-Trp; Xaa at residues 11 and 19 is Glu or gamma-carboxy-Glu.

<220>

<221> SITE

<222> (12)..(16)

<223> Xaa at residues 12 and 16 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys; Xaa at residue 13 is Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<400> 186

Xaa Asn Cys Cys Ser Ile Xaa Gly Cys Xaa Xaa Xaa Xaa Gly Asp Xaa
1 5 10 15

Cys Ser Xaa Val Arg
20

<210> 187

<211> 16

<212> PRT

<213> Conus catus

<220>

<221> SITE

<222> (6)..(13)

<223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro; Xaa at residue 11 is Glu or gamma-carboxy-Glu.

<400> 187

Gly Cys Cys Ser Asn Xaa Val Cys His Leu Xaa His Xaa Asn Ala Cys
1 5 10 15

<210> 188

<211> 17

<212> PRT

<213> Conus catus

<220>

<221> SITE

<222> (6)..(13)

<223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro; Xaa at residue 9 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr.

<400> 188

Gly Cys Cys Ser Asn Xaa Ile Cys Xaa Phe Asn Asn Xaa Arg Ile Cys
1 5 10 15

Arg

<210> 189

<211> 17

<212> PRT

<213> Conus episcopatus

<220>

<221> SITE

<222> (1)..(14)
 <223> Xaa at residues 1 and 14 is Glu or
 gamma-carboxy-Glu; Xaa at residues 6, 7 and 13 is
 Pro or hydroxy-Pro; Xaa at residue 10 is Trp (D or
 L) or halo-Trp.

<220>
 <221> SITE
 <222> (11)
 <223> Xaa at residue 11 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 189
 Xaa Cys Cys Ser Gln Xaa Xaa Cys Arg Xaa Xaa His Xaa Xaa Leu Cys
 1 5 10 15
 Ser

<210> 190
 <211> 16
 <212> PRT
 <213> Conus geographus

<220>
 <221> SITE
 <222> (6)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro.

<400> 190
 Gly Cys Cys Ser His Xaa Ala Cys Ala Gly Asn Asn Gln His Ile Cys
 1 5 10 15

<210> 191
 <211> 18
 <212> PRT
 <213> Conus geographus

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 191
 Gly Cys Cys Ala Val Xaa Ser Cys Arg Leu Arg Asn Xaa Asp Leu Cys
 1 5 10 15
 Gly Gly

<210> 192
 <211> 16
 <212> PRT
 <213> Conus imperialis

<220>
 <221> NP_BIND
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 192

Gly Cys Cys Ser His Xaa Ala Cys Asn Val Asn Asn Xaa His Ile Cys
 1 5 10 15

<210> 193
 <211> 20
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (2)..(10)
 <223> Xaa at residues 2, 7, 9 and 10 is Pro or
 hydroxy-Pro; Xaa at residues 3 and 4 is Glu or
 gamma-carboxy-Glu.

<400> 193
 Thr Xaa Xaa Xaa Cys Cys Xaa Asn Xaa Xaa Cys Phe Ala Thr Asn Ser
 1 5 10 15

Asp Ile Cys Gly
 20

<210> 194
 <211> 17
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (7)..(12)
 <223> Xaa at residue 7 is Pro or hydroxy-Pro; Xaa at
 residue 12 is Lys, N-methyl-Lys, N,N-dimethyl-Lys
 or N,N,N-trimethyl-Lys.

<400> 194
 Asp Ala Cys Cys Ser Asp Xaa Arg Cys Ser Gly Xaa His Gln Asp Leu
 1 5 10 15

Cys

<210> 195
 <211> 17
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (1)..(7)
 <223> Xaa at residue 1 is Glu or gamma-carboxy-Glu; Xaa
 at residue 7 is Pro or hydroxy-Pro.

<400> 195
 Xaa Asp Cys Cys Ser Asp Xaa Arg Cys Ser Val Gly His Gln Asp Leu
 1 5 10 15

Cys

<210> 196

<211> 16
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (6)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro.

<400> 196
 Gly Cys Cys Ser His Xaa Ala Cys Ala Gly Ser Asn Ala His Ile Cys
 1 5 10 15

<210> 197
 <211> 17
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (1)..(7)
 <223> Xaa at residue 1 is Glu or gamma-carboxy-Glu; Xaa
 at residue 7 is Pro or hydroxy-Pro.

<400> 197
 Xaa Asp Cys Cys Ser Asp Xaa Arg Cys Ser Val Gly His Gln Asp Met
 1 5 10 15

Cys

<210> 198
 <211> 16
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 198
 Gly Cys Cys Ser His Xaa Ala Cys Ala Gly Asn Asn Xaa His Ile Cys
 1 5 10 15

<210> 199
 <211> 17
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (6)..(14)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 14 is Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 199
 Gly Cys Cys Gly Asn Xaa Ser Cys Ser Ile His Ile Xaa Xaa Val Cys
 1 5 10 15

Asn

<210> 200
 <211> 21
 <212> PRT
 <213> Conus lividus

<220>
 <221> SITE
 <222> (4)..(5)
 <223> Xaa at residues 4 and 5 is Glu or
 gamma-carboxy-Glu.

<400> 200
 Thr Asp Ser Xaa Xaa Cys Cys Leu Asp Ser Arg Cys Ala Gly Gln His
 1 5 10 15

Gln Asp Leu Cys Gly
 20

<210> 201
 <211> 17
 <212> PRT
 <213> Conus marmoreus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residues 6 and 7 is Pro or hydroxy-Pro; Xaa
 at residues 9 and 15 is Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 201
 Gly Cys Cys Ser Asn Xaa Xaa Cys Xaa Ala Asn Asn Gln Ala Xaa Cys
 1 5 10 15

Asn

<210> 202
 <211> 16
 <212> PRT
 <213> Conus marmoreus

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 202
 Gly Cys Cys Ser His Xaa Ala Cys Ser Val Asn Asn Xaa Asp Ile Cys
 1 5 10 15

<210> 203
 <211> 18
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (2)..(15)
 <223> Xaa at residues 2 and 12 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys; Xaa at
 residue 14 is Pro or hydroxy-Pro.

<220>
 <221> SITE
 <222> (16)
 <223> Xaa at residue 16 is Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 203
 Gly Xaa Cys Cys Ile Asn Asp Ala Cys Arg Ser Xaa His Xaa Gln Xaa
 1 5 10 15

Cys Ser

<210> 204
 <211> 17
 <212> PRT
 <213> Conus musicus

<220>
 <221> SITE
 <222> (4)..(15)
 <223> Xaa at residues 4 and 15 is Tyr, nor-Tyr,
 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr or nitro-Tyr; Xaa at residue 13 is
 Pro or hydroxy-Pro.

<400> 204
 Gly Cys Cys Xaa Asn Ile Ala Cys Arg Ile Asn Asn Xaa Arg Xaa Cys
 1 5 10 15

Arg

<210> 205
 <211> 17
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residues 12 and 15 is Tyr, nor-Tyr,
 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr or nitro-Tyr.

<220>
 <221> SITE
 <222> (14)
 <223> Xaa at residue 14 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 205
 Gly Cys Cys Ser His Xaa Val Cys Arg Phe Asn Xaa Xaa Xaa Xaa Cys

1 5 10 15
 Gly

<210> 206
 <211> 18
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (2)..(15)
 <223> Xaa at residue 2 is Glu or gamma-carboxy-Glu; Xaa
 at residues 7, 8 and 14 is Pro or hydroxy-Pro; Xaa
 at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>
 <221> SITE
 <222> (2)..(15)
 <223> nitro-Tyr

<400> 206
 Asp Xaa Cys Cys Ala Ser Xaa Xaa Cys Arg Leu Asn Asn Xaa Xaa Val
 1 5 10 15

Cys His

<210> 207
 <211> 19
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (6)..(18)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residue 9 is Trp (D or L) or halo-Trp; Xaa at
 residues 14 and 18 is Glu or gamma-carboxy-Glu.

<220>
 <221> SITE
 <222> (15)
 <223> Xaa at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 207
 Gly Cys Cys Ser Asn Xaa Val Cys Xaa Gln Asn Asn Ala Xaa Xaa Cys
 1 5 10 15

Arg Xaa Ser

<210> 208
 <211> 16
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residues 6 and 7 is Pro or hydroxy-Pro; Xaa
 at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

 <400> 208
 Gly Cys Cys Ser His Xaa Xaa Cys Ala Gln Asn Asn Gln Asp Xaa Cys
 1 5 10 15

<210> 209
 <211> 19
 <212> PRT
 <213> Conus obscurus

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residues 14 and 18 is Glu or gamma-carboxy-Glu;
 Xaa at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or

<220>
 <221> SITE
 <222> (6)..(15)
 <223> nitro-Tyr.

<400> 209
 Gly Cys Cys Ser His Xaa Ala Cys Ser Gly Asn Asn Arg Xaa Xaa Cys
 1 5 10 15

Arg Xaa Ser

<210> 210
 <211> 18
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (2)..(15)
 <223> Xaa at residues 2, 7 and 14 is Pro or hydroxy-Pro;
 Xaa at residue 6 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr; Xaa at residue 15 is Glu or

<220>
 <221> SITE
 <222> (2)..(15)
 <223> gamma-carboxy-Glu

<400> 210
 Asp Xaa Cys Cys Ser Xaa Xaa Asp Cys Gly Ala Asn His Xaa Xaa Ile
 1 5 10 15

Cys Gly

<210> 211
 <211> 17
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (1)..(14)
 <223> Xaa at residues 1 and 14 is Glu or
 gamma-carboxy-Glu; Xaa at residues 6, 7 and 13 is
 Pro or hydroxy-Pro; Xaa at residue 10 is Trp (D or
 L) or halo-Trp.

<220>
 <221> SITE
 <222> (11)
 <223> Xaa at residue 11 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 211
 Xaa Cys Cys Ser Gln Xaa Xaa Cys Arg Xaa Xaa His Xaa Xaa Leu Cys
 1 5 10 15

Ser

<210> 212
 <211> 16
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 212
 Gly Cys Cys Ser His Xaa Ala Cys Ala Gly Asn Asn Xaa His Ile Cys
 1 5 10 15

<210> 213
 <211> 16
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (6)..(15)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 15 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 213
 Gly Cys Cys Ser Asp Xaa Ser Cys Asn Val Asn Asn Xaa Asp Xaa Cys
 1 5 10 15

<210> 214
 <211> 18
 <212> PRT
 <213> Conus omaria

<220>
 <221> SITE
 <222> (1)..(7)
 <223> Xaa at residues 1 and 2 is Glu or
 gamma-carboxy-Glu; Xaa at residue 7 is Pro or
 hydroxy-Pro.

<400> 214
 Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Ser Val Gly His Gln Asp Met
 1 5 10 15

Cys Arg

<210> 215
 <211> 17
 <212> PRT
 <213> Conus purpurascens

<220>
 <221> SITE
 <222> (7)..(15)
 <223> Xaa at residue 7 is Pro or hydroxy-Pro; Xaa at
 residue 15 is Glu or gamma-carboxy-Glu.

<400> 215
 Gly Gly Cys Cys Ser Asn Xaa Ala Cys Leu Val Asn His Leu Xaa Met
 1 5 10 15

Cys

<210> 216
 <211> 18
 <212> PRT
 <213> Conus purpurascens

<220>
 <221> SITE
 <222> (3)..(15)
 <223> Xaa at residues 3, 8 and 15 is Pro or hydroxy-Pro.

<400> 216
 Arg Asp Xaa Cys Cys Phe Asn Xaa Ala Cys Asn Val Asn Asn Xaa Gln
 1 5 10 15

Ile Cys

<210> 217
 <211> 21
 <212> PRT
 <213> Conus purpurascens

<220>
 <221> SITE
 <222> (5)..(8)
 <223> Xaa at residue 5 is Pro or hydroxy-Pro; Xaa at
 residue 8 is Trp (D or L) or halo-Trp.

<400> 217

Cys Cys Ser Asp Xaa Ser Cys Xaa Arg Leu His Ser Leu Ala Cys Thr
 1 5 10 15

Gly Ile Val Asn Arg
 20

<210> 218
 <211> 16
 <212> PRT
 <213> Conus purpurascens

<220>
 <221> SITE
 <222> (5)
 <223> Xaa at residue 5 is Pro or hydroxy-Pro.

<400> 218
 Cys Cys Thr Asn Xaa Ala Cys Leu Val Asn Asn Ile Arg Phe Cys Gly
 1 5 10 15

<210> 219
 <211> 18
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (2)..(7)
 <223> Xaa at residue 2 is Glu or gamma-carboxy-Glu; Xaa
 at residue 7 is Pro or hydroxy-Pro.

<400> 219
 Asp Xaa Cys Cys Ser Asp Xaa Arg Cys His Gly Asn Asn Arg Asp His
 1 5 10 15

Cys Ala

<210> 220
 <211> 17
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 220
 Asp Cys Cys Ser His Xaa Leu Cys Arg Leu Phe Val Xaa Gly Leu Cys
 1 5 10 15

Ile

<210> 221
 <211> 17
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
 Xaa at residue 9 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<220>
 <221> SITE
 <222> (12)
 <223> Xaa at residue 12 is Tyr, nor-Tyr, mono-halo-Tyr,
 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or
 nitro-Tyr.

<400> 221
 Gly Cys Cys Ser His Xaa Val Cys Xaa Val Arg Xaa Xaa Asp Leu Cys
 1 5 10 15

Arg

<210> 222
 <211> 16
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (6)..(13)
 <223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 222
 Gly Cys Cys Ser His Xaa Ala Cys Asn Val Asn Asn Xaa His Ile Cys
 1 5 10 15

<210> 223
 <211> 16
 <212> PRT
 <213> Conus regius

<220>
 <221> SITE
 <222> (6)..(12)
 <223> Xaa at residue 6 is Pro or hydroxy-Pro; Xaa at
 residue 12 is Tyr, nor-Tyr, nor-Tyr,
 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
 O-phospho-Tyr or nitro-Tyr.

<220>
 <221> SITE
 <222> (9)
 <223> Xaa at residue 9 is Lys, N-methyl-Lys,
 N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 223
 Gly Cys Cys Ser His Xaa Val Cys Xaa Val Arg Xaa Ser Asp Met Cys
 1 5 10 15

<210> 224
 <211> 17
 <212> PRT

<213> Conus stercusmuscarum

<220>

<221> SITE

<222> (7)..(14)

<223> Xaa at residues 7 and 14 is Pro or hydroxy-Pro;
Xaa at residue 10 is Lys, N-methyl-Lys,
N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 224

Gly Gly Cys Cys Ser His Xaa Ala Cys Xaa Val His Phe Xaa His Ser
1 5 10 15

Cys

<210> 225

<211> 20

<212> PRT

<213> Conus stercusmuscarum

<220>

<221> SITE

<222> (6)..(14)

<223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro;
Xaa at residue 14 is Glu or gamma-carboxy-Glu.

<400> 225

Val Cys Cys Ser Asn Xaa Val Cys His Val Asp His Xaa Xaa Leu Cys
1 5 10 15

Arg Arg Arg Arg
20

<210> 226

<211> 17

<212> PRT

<213> Conus striatus

<220>

<221> SITE

<222> (6)..(13)

<223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 226

Gly Cys Cys Ser His Xaa Val Cys Asn Leu Ser Asn Xaa Gln Ile Cys
1 5 10 15

Arg

<210> 227

<211> 18

<212> PRT

<213> Conus textile

<220>

<221> SITE

<222> (1)..(15)

<223> Xaa at residue 1 is Gln or pyro-Glu; Xaa at
residues 2 and 15 is Glu or gamma-carboxy-Glu; Xaa

at residues 7 and 14 is Pro or hydroxy-Pro.

<400> 227

Xaa Xaa Cys Cys Ser His Xaa Ala Cys Asn Val Asp His Xaa Xaa Ile
1 5 10 15

Cys Arg

<210> 228

<211> 17

<212> PRT

<213> Conus tulipa

<220>

<221> SITE

<222> (6)

<223> Xaa at residue 6 is Pro or hydroxy-Pro.

<400> 228

Gly Cys Cys Ser Asn Xaa Ala Cys Leu Val Asn His Ile Arg Phe Cys
1 5 10 15

Gly

<210> 229

<211> 17

<212> PRT

<213> Conus virgo

<220>

<221> SITE

<222> (6)..(13)

<223> Xaa at residues 6 and 13 is Pro or hydroxy-Pro.

<400> 229

Asp Cys Cys Asp Asp Xaa Ala Cys Thr Val Asn Asn Xaa Gly Leu Cys
1 5 10 15

Thr

<210> 230

<211> 20

<212> PRT

<213> Conus textile

<220>

<221> SITE

<222> (6)..(13)

<223> Xaa at residues 6, 7 and 13 is Pro or hydroxy-Pro;
Xaa at residue 11 is Lys, N-methyl-Lys,
N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 230

Gly Cys Cys Ser Asn Xaa Xaa Cys Ile Ala Xaa Asn Xaa His Met Cys
1 5 10 15

Gly Gly Arg Arg
20

<210> 231
 <211> 18
 <212> PRT
 <213> *Conus geographus*

<220>
 <221> SITE
 <222> (5)..(9)
 <223> Xaa at residue 5 is Pro or hydroxy-Pro; Xaa at residue 8 is Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr or nitro-Tyr; Xaa at residue 9 is Glu or

<220>
 <221> SITE
 <222> (9)..(14)
 <223> gamma-carboxy-Glu; Xaa at residues 10, 11, 12 and 14 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 231
 Cys Cys Thr Ile Xaa Ser Cys Xaa Xaa Xaa Xaa Xaa Ile Xaa Ala Cys
 1 5 10 15
 Val Phe

<210> 232
 <211> 18
 <212> PRT
 <213> *Conus regius*

<220>
 <221> SITE
 <222> (6)..(16)
 <223> Xaa at residues 6 and 16 is Pro or hydroxy-Pro; Xaa at residue 13 is Lys, N-methyl-Lys, N,N-dimethyl-Lys or N,N,N-trimethyl-Lys.

<400> 232
 Gly Cys Cys Gly Asn Xaa Ala Cys Ser Gly Ser Ser Xaa Asp Ala Xaa
 1 5 10 15
 Ser Cys

<210> 233
 <211> 108
 <212> DNA
 <213> *Conus imperialis*

<220>
 <221> CDS
 <222> (1)..(105)

<400> 233
 tct gat gga aag agt gcc gcg gcc aaa gcc aaa ccg tct cac ctg acg 48
 Ser Asp Gly Lys Ser Ala Ala Ala Lys Ala Lys Pro Ser His Leu Thr
 1 5 10 15
 gct cca ttc atc agg gac gaa tgc tgt tcc gat tct cgc tgt ggc aag 96

Ala Pro Phe Ile Arg Asp Glu Cys Cys Ser Asp Ser Arg Cys Gly Lys
 20 25 30

aac tgt ctt tga
 Asn Cys Leu
 35

108

<210> 234
 <211> 35
 <212> PRT
 <213> Conus imperialis

<400> 234
 Ser Asp Gly Lys Ser Ala Ala Ala Lys Ala Lys Pro Ser His Leu Thr
 1 5 10 15

Ala Pro Phe Ile Arg Asp Glu Cys Cys Ser Asp Ser Arg Cys Gly Lys
 20 25 30

Asn Cys Leu
 35

<210> 235
 <211> 108
 <212> DNA
 <213> Conus imperialis

<220>
 <221> CDS
 <222> (1)..(105)

<400> 235
 ttt gat gga agg aat gcc cca gcc gac gac aaa gcg tct gac ctg atc 48
 Phe Asp Gly Arg Asn Ala Pro Ala Asp Asp Lys Ala Ser Asp Leu Ile
 1 5 10 15

gct caa atc gtc agg aga gca tgc tgt tcc gat cgt cgc tgt aga tgg 96
 Ala Gln Ile Val Arg Arg Ala Cys Cys Ser Asp Arg Arg Cys Arg Trp
 20 25 30

agg tgt ggt tga
 Arg Cys Gly
 35 108

<210> 236
 <211> 35
 <212> PRT
 <213> Conus imperialis

<400> 236
 Phe Asp Gly Arg Asn Ala Pro Ala Asp Asp Lys Ala Ser Asp Leu Ile
 1 5 10 15

Ala Gln Ile Val Arg Arg Ala Cys Cys Ser Asp Arg Arg Cys Arg Trp
 20 25 30

Arg Cys Gly
 35

<210> 237

<211> 145
 <212> DNA
 <213> Conus regius

<220>
 <221> CDS
 <222> (1)..(105)

<400> 237
 tct gat gga agg aat gcc gca gcc gac gcc aga gcg tct ccc cgg atc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asp Ala Arg Ala Ser Pro Arg Ile
 1 5 10 15
 gct ctt ttc ctc agg ttc aca tgc tgt agg aga ggt acc tgt tcc cag 96
 Ala Leu Phe Leu Arg Phe Thr Cys Cys Arg Arg Gly Thr Cys Ser Gln
 20 25 30
 cac tgt ggt tgaagacact gctgctccag gaccctctga accacgacgt 145
 His Cys Gly
 35

<210> 238
 <211> 35
 <212> PRT
 <213> Conus regius

<400> 238
 Ser Asp Gly Arg Asn Ala Ala Ala Asp Ala Arg Ala Ser Pro Arg Ile
 1 5 10 15
 Ala Leu Phe Leu Arg Phe Thr Cys Cys Arg Arg Gly Thr Cys Ser Gln
 20 25 30
 His Cys Gly
 35

<210> 239
 <211> 145
 <212> DNA
 <213> Conus regius

<220>
 <221> CDS
 <222> (1)..(105)

<400> 239
 tct aat gga agg aat gcc gca gcc gac gcc aaa gcg tct caa cgg atc 48
 Ser Asn Gly Arg Asn Ala Ala Ala Asp Ala Lys Ala Ser Gln Arg Ile
 1 5 10 15
 gct cca ttc ctc agg gac tat tgc tgt agg aga cat gcc tgt acg ttg 96
 Ala Pro Phe Leu Arg Asp Tyr Cys Cys Arg Arg His Ala Cys Thr Leu
 20 25 30
 att tgt ggt tgaagacgct gctgctccag gaccctctga accacgacgt 145
 Ile Cys Gly
 35

<210> 240
 <211> 35
 <212> PRT

<213> Conus regius

<400> 240

Ser Asn Gly Arg Asn Ala Ala Ala Asp Ala Lys Ala Ser Gln Arg Ile
1 5 10 15

Ala Pro Phe Leu Arg Asp Tyr Cys Cys Arg Arg His Ala Cys Thr Leu
20 25 30

Ile Cys Gly
35

<210> 241

<211> 145

<212> DNA

<213> Conus regius

<220>

<221> CDS

<222> (1)..(105)

<400> 241

tct aat gga agg aat gcc gca gcc gac gcc aaa gcg tct caa cgg atc 48
Ser Asn Gly Arg Asn Ala Ala Ala Asp Ala Lys Ala Ser Gln Arg Ile
1 5 10 15

gct cca ttc ctc agg gac tat tgc tgt agg aga cct ccc tgt acg ttg 96
Ala Pro Phe Leu Arg Asp Tyr Cys Cys Arg Arg Pro Pro Cys Thr Leu
20 25 30

att tgt ggt tgaagacgct gctgctccag gaccctctga accacgacgt 145
Ile Cys Gly
35

<210> 242

<211> 35

<212> PRT

<213> Conus regius

<400> 242

Ser Asn Gly Arg Asn Ala Ala Ala Asp Ala Lys Ala Ser Gln Arg Ile
1 5 10 15

Ala Pro Phe Leu Arg Asp Tyr Cys Cys Arg Arg Pro Pro Cys Thr Leu
20 25 30

Ile Cys Gly
35

<210> 243

<211> 136

<212> DNA

<213> Conus regius

<220>

<221> CDS

<222> (1)..(96)

<400> 243

tct aat aaa agg aag aat gcc gca atg ctt gac atg atc gct caa cac 48
Ser Asn Lys Arg Lys Asn Ala Ala Met Leu Asp Met Ile Ala Gln His

1 5 10 15
gcc ata agg ggt tgc tgt tcc gat cct cgc tgt aga tat aga tgt cgt 96
Ala Ile Arg Gly Cys Cys Ser Asp Pro Arg Cys Arg Tyr Arg Cys Arg
20 25 30
tgaagacgct gctgctccag gaccctctga accacgacgt 136

<210> 244
<211> 32
<212> PRT
<213> Conus regius

<400> 244
Ser Asn Lys Arg Lys Asn Ala Ala Met Leu Asp Met Ile Ala Gln His
1 5 10 15
Ala Ile Arg Gly Cys Cys Ser Asp Pro Arg Cys Arg Tyr Arg Cys Arg
20 25 30

<210> 245
<211> 145
<212> DNA
<213> Conus regius

<220>
<221> CDS
<222> (1)..(105)

<400> 245
ttt aat gga agg agt gcc gca gcc gac caa aat gcg cct ggc ctg atc 48
Phe Asn Gly Arg Ser Ala Ala Ala Asp Gln Asn Ala Pro Gly Leu Ile
1 5 10 15
gct caa gtc gtc aga gga ggg tgc tgt tcc gat ccc cgc tgc gcc tgg 96
Ala Gln Val Val Arg Gly Gly Cys Cys Ser Asp Pro Arg Cys Ala Trp
20 25 30
aga tgt ggt tgaagacgctt gctgctccag gaccctctga accacgacgt 145
Arg Cys Gly
35

<210> 246
<211> 35
<212> PRT
<213> Conus regius

<400> 246
Phe Asn Gly Arg Ser Ala Ala Ala Asp Gln Asn Ala Pro Gly Leu Ile
1 5 10 15
Ala Gln Val Val Arg Gly Gly Cys Cys Ser Asp Pro Arg Cys Ala Trp
20 25 30
Arg Cys Gly
35

<210> 247
<211> 145
<212> DNA

<213> Conus regius

<220>

<221> CDS

<222> (1)..(105)

<400> 247

ttt gat gga agg aat gcc gca gcc gac gcc aaa gtg att aac acg gtc	48
Phe Asp Gly Arg Asn Ala Ala Ala Asp Ala Lys Val Ile Asn Thr Val	
1 5 10 15	

gct cga atc gcc tgg gat ata tgc tgt tcc gaa cct gac tgt aac cat	96
Ala Arg Ile Ala Trp Asp Ile Cys Cys Ser Glu Pro Asp Cys Asn His	
20 25 30	

aaa tgt gtt tgaagacgct tctgctccag gaccctctga accacgacgt	145
Lys Cys Val	
35	

<210> 248

<211> 35

<212> PRT

<213> Conus regius

<400> 248

Phe Asp Gly Arg Asn Ala Ala Ala Asp Ala Lys Val Ile Asn Thr Val	
1 5 10 15	

Ala Arg Ile Ala Trp Asp Ile Cys Cys Ser Glu Pro Asp Cys Asn His	
20 25 30	

Lys Cys Val	
35	

<210> 249

<211> 136

<212> DNA

<213> Conus regius

<220>

<221> CDS

<222> (1)..(96)

<400> 249

tct aat aaa agg aag aat gcc gca atg ctt gac atg atc gct caa cac	48
Ser Asn Lys Arg Lys Asn Ala Ala Met Leu Asp Met Ile Ala Gln His	
1 5 10 15	

gcc ata agg ggt tgc tgt tcc gat cct cgc tgt aaa cat cag tgt ggt	96
Ala Ile Arg Gly Cys Cys Ser Asp Pro Arg Cys Lys His Gln Cys Gly	
20 25 30	

tgaagacgct gctgctccag gaccctctga accacgacgt	136
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<210> 250

<211> 32

<212> PRT

<213> Conus regius

<400> 250

Ser Asn Lys Arg Lys Asn Ala Ala Met Leu Asp Met Ile Ala Gln His	
---	--

1 5 10 15

Ala Ile Arg Gly Cys Cys Ser Asp Pro Arg Cys Lys His Gln Cys Gly
 20 25 30

<210> 251
 <211> 136
 <212> DNA
 <213> Conus musicus

<220>
 <221> CDS
 <222> (1)..(105)

<400> 251
 atc aag aat aca gca gcc agc aac aaa gcg tct agc ctg gtg gct ctt 48
 Ile Lys Asn Thr Ala Ala Ser Asn Lys Ala Ser Ser Leu Val Ala Leu
 1 5 10 15

gtt gtc agg gga tgc tgt tac aat cct gtc tgc aag aaa tat tat tgt 96
 Val Val Arg Gly Cys Cys Tyr Asn Pro Val Cys Lys Lys Tyr Tyr Cys
 20 25 30

tgg aaa ggc tgatgctcca ggaccctctg aaccacgacg t 136
 Trp Lys Gly
 35

<210> 252
 <211> 35
 <212> PRT
 <213> Conus musicus

<400> 252
 Ile Lys Asn Thr Ala Ala Ser Asn Lys Ala Ser Ser Leu Val Ala Leu
 1 5 10 15

Val Val Arg Gly Cys Cys Tyr Asn Pro Val Cys Lys Lys Tyr Tyr Cys
 20 25 30

Trp Lys Gly
 35

<210> 253
 <211> 148
 <212> DNA
 <213> Conus purpurascens

<220>
 <221> CDS
 <222> (1)..(117)

<400> 253
 tct gaa ggc agg aat gct gaa gcc atc gac aac gcc tta gac cag agg 48
 Ser Glu Gly Arg Asn Ala Glu Ala Ile Asp Asn Ala Leu Asp Gln Arg
 1 5 10 15

gat cca aag cga cag gag ccg ggg tgc tgt agg cat cct gcc tgt ggg 96
 Asp Pro Lys Arg Gln Glu Pro Gly Cys Cys Arg His Pro Ala Cys Gly
 20 25 30

aag aac aga tgt gga aga cgc tgatgctcca ggaccctctg aaccacgacg t 148

Lys Asn Arg Cys Gly Arg Arg
35

<210> 254
<211> 39
<212> PRT
<213> *Conus purpurascens*

<400> 254
Ser Glu Gly Arg Asn Ala Glu Ala Ile Asp Asn Ala Leu Asp Gln Arg
1 5 10 15
Asp Pro Lys Arg Gln Glu Pro Gly Cys Cys Arg His Pro Ala Cys Gly
20 25 30

Lys Asn Arg Cys Gly Arg Arg
35

<210> 255
<211> 156
<212> DNA
<213> *Conus musicus*

<220>
<221> CDS
<222> (1)..(102)

<400> 255
tct gat ggc agg aat att gca gtc gac gac aga tgg tct ttc tat acg 48
Ser Asp Gly Arg Asn Ile Ala Val Asp Asp Arg Trp Ser Phe Tyr Thr
1 5 10 15
ctc ttc cat gct act tgc tgt gcc gat cct gac tgt aga ttc cgg ccc 96
Leu Phe His Ala Thr Cys Cys Ala Asp Pro Asp Cys Arg Phe Arg Pro
20 25 30
ggg tgt tgatctttgt tcttcaaaga cgctgctggc ccaggaccct ctgaaccacg 152
Gly Cys
acgt 156

<210> 256
<211> 34
<212> PRT
<213> *Conus musicus*

<400> 256
Ser Asp Gly Arg Asn Ile Ala Val Asp Asp Arg Trp Ser Phe Tyr Thr
1 5 10 15
Leu Phe His Ala Thr Cys Cys Ala Asp Pro Asp Cys Arg Phe Arg Pro
20 25 30

Gly Cys

<210> 257
<211> 142
<212> DNA
<213> *Conus musicus*

<220>
 <221> CDS
 <222> (1)..(102)

<400> 257

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atc aag aat act gca gcc agc aac aaa gcg cct agc ctg gtg gct att   48
Ile Lys Asn Thr Ala Ala Ser Asn Lys Ala Pro Ser Leu Val Ala Ile
  1             5             10             15

gcc gtc agg gga tgc tgt tac aat cct tcc tgt tgg ccg aaa aca tat   96
Ala Val Arg Gly Cys Cys Tyr Asn Pro Ser Cys Trp Pro Lys Thr Tyr
             20             25             30

tgt agt tggaaaggct gatgctccag gaccctctga accacgacgt   142
Cys Ser

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<210> 258
 <211> 34
 <212> PRT
 <213> Conus musicus

<400> 258

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Ile Lys Asn Thr Ala Ala Ser Asn Lys Ala Pro Ser Leu Val Ala Ile
  1             5             10             15

Ala Val Arg Gly Cys Cys Tyr Asn Pro Ser Cys Trp Pro Lys Thr Tyr
             20             25             30

Cys Ser

```

<210> 259
 <211> 161
 <212> DNA
 <213> Conus musicus

<220>
 <221> CDS
 <222> (1)..(108)

<400> 259

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tct gat agc agg aat gtc gca atc gag gac aga gtg tct gac ctg cac   48
Ser Asp Ser Arg Asn Val Ala Ile Glu Asp Arg Val Ser Asp Leu His
  1             5             10             15

tct atg ttc ttc gat gtt tct tgc tgt agc aat cct acc tgt aaa gaa   96
Ser Met Phe Phe Asp Val Ser Cys Cys Ser Asn Pro Thr Cys Lys Glu
             20             25             30

acg tat ggt tgt tgatcggttg ttttgaagac gctgatgctc caggaccctc   148
Thr Tyr Gly Cys
             35

tgaaccacga cgt   161

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<210> 260
 <211> 36
 <212> PRT
 <213> Conus musicus

<400> 260

Ser Asp Ser Arg Asn Val Ala Ile Glu Asp Arg Val Ser Asp Leu His
 1 5 10 15

Ser Met Phe Phe Asp Val Ser Cys Cys Ser Asn Pro Thr Cys Lys Glu
 20 25 30

Thr Tyr Gly Cys
 35

<210> 261

<211> 156

<212> DNA

<213> Conus musicus

<220>

<221> CDS

<222> (1)..(102)

<400> 261

tct gtt ggc agg aat att gca gtc gac gac aga ggg att ttc tct acg 48
 Ser Val Gly Arg Asn Ile Ala Val Asp Asp Arg Gly Ile Phe Ser Thr
 1 5 10 15

ctc ttc cat gct cat tgc tgt gcc aat ccc atc tgt aaa aac acg ccc 96
 Leu Phe His Ala His Cys Cys Ala Asn Pro Ile Cys Lys Asn Thr Pro
 20 25 30

ggc tgt tgatctttgt tcttcaaaga cgctgctggc ccaggaccct ctgaaccacg 152
 Gly Cys

acgt 156

<210> 262

<211> 34

<212> PRT

<213> Conus musicus

<400> 262

Ser Val Gly Arg Asn Ile Ala Val Asp Asp Arg Gly Ile Phe Ser Thr
 1 5 10 15

Leu Phe His Ala His Cys Cys Ala Asn Pro Ile Cys Lys Asn Thr Pro
 20 25 30

Gly Cys

<210> 263

<211> 161

<212> DNA

<213> Conus musicus

<220>

<221> CDS

<222> (1)..(108)

<400> 263

tcc gat ggc agg aat gtc gca atc gac gac aga gtg tct gac ctg cac 48
 Ser Asp Gly Arg Asn Val Ala Ile Asp Asp Arg Val Ser Asp Leu His
 1 5 10 15

tct atg ttc ttc gat att gct tgc tgt aac aat cct acc tgt aaa gaa 96
Ser Met Phe Phe Asp Ile Ala Cys Cys Asn Asn Pro Thr Cys Lys Glu
20 25 30

acg tat ggt tgt tgatcgttgg ttttgaagac gctgatgctc caggaccctc 148
Thr Tyr Gly Cys
35

tgaaccacga cgt 161

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<210> 264
<211> 36
<212> PRT
<213> Conus musicus
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<400> 264
Ser Asp Gly Arg Asn Val Ala Ile Asp Asp Arg Val Ser Asp Leu His
1 5 10 15

Ser Met Phe Phe Asp Ile Ala Cys Cys Asn Asn Pro Thr Cys Lys Glu
20 25 30

Thr Tyr Gly Cys
35

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<210> 265
<211> 161
<212> DNA
<213> Conus musicus
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<220>
<221> CDS
<222> (1) .. (108)
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<400> 265
tct gat ggc agg aat gtc gca atc gag gac aga gtg tct gac ctg ctc 48
Ser Asp Gly Arg Asn Val Ala Ile Glu Asp Arg Val Ser Asp Leu Leu
  1             5             10             15
```

tct atg ctc ttc gat gtt gct tgc tgt agc aat cct gtc tgt aaa gaa 96
Ser Met Leu Phe Asp Val Ala Cys Cys Ser Asn Pro Val Cys Lys Glu
20 25 30

acg tat ggt tgt tgatcgttgg ttttgaagac gctgatgctc caggaccctc 148
Thr Tyr Gly Cys
35

tgaaccacga cgt 161

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<210> 266
<211> 36
<212> PRT
<213> Conus musicus
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<400> 266
Ser Asp Gly Arg Asn Val Ala Ile Glu Asp Arg Val Ser Asp Leu Leu
1 5 10 15

Ser Met Leu Phe Asp Val Ala Cys Cys Ser Asn Pro Val Cys Lys Glu
20 25 30

Thr Tyr Gly Cys
35

<210> 267
<211> 154
<212> DNA
<213> Conus betulinus

<220>
<221> CDS
<222> (1)..(123)

<400> 267
tat gat ggc agg aat gct gcc gcc gac gac aaa gct ttt gac ctg ctg 48
Tyr Asp Gly Arg Asn Ala Ala Ala Asp Asp Lys Ala Phe Asp Leu Leu
1 5 10 15

gct atg acc ata agg gga gga tgc tgt tcc tat cct ccc tgt atc gcg 96
Ala Met Thr Ile Arg Gly Gly Cys Cys Ser Tyr Pro Pro Cys Ile Ala
20 25 30

agt aat cct aaa tgt ggt gga aga cgc tgatgctcca ggaccctctg 143
Ser Asn Pro Lys Cys Gly Gly Arg Arg
35 40

aaccacaacg t 154

<210> 268
<211> 41
<212> PRT
<213> Conus betulinus

<400> 268
Tyr Asp Gly Arg Asn Ala Ala Ala Asp Asp Lys Ala Phe Asp Leu Leu
1 5 10 15

Ala Met Thr Ile Arg Gly Gly Cys Cys Ser Tyr Pro Pro Cys Ile Ala
20 25 30

Ser Asn Pro Lys Cys Gly Gly Arg Arg
35 40

<210> 269
<211> 151
<212> DNA
<213> Conus lividus

<220>
<221> CDS
<222> (1)..(111)

<400> 269
ttt gat ggc agg aat gct gca ggc aac gcc aaa atg tcc gcc ctg atg 48
Phe Asp Gly Arg Asn Ala Ala Gly Asn Ala Lys Met Ser Ala Leu Met
1 5 10 15

gcc ctg acc atc agg gga tgc tgt tcc cat cct gtc tgt agc gcg atg 96
Ala Leu Thr Ile Arg Gly Cys Cys Ser His Pro Val Cys Ser Ala Met
20 25 30

agt cca atc tgt ggc tgaagacgct gatgccccag gaccctctga accacgacgt 151

Ser Pro Ile Cys Gly
35

<210> 270
<211> 37
<212> PRT
<213> Conus lividus

<400> 270
Phe Asp Gly Arg Asn Ala Ala Gly Asn Ala Lys Met Ser Ala Leu Met
1 5 10 15
Ala Leu Thr Ile Arg Gly Cys Cys Ser His Pro Val Cys Ser Ala Met
20 25 30
Ser Pro Ile Cys Gly
35

<210> 271
<211> 196
<212> DNA
<213> Conus musicus

<220>
<221> CDS
<222> (1)...(165)

<400> 271
atc aag aat gct gca gct gac gac aaa gca tct gac ctg ctc tct cag 48
Ile Lys Asn Ala Ala Ala Asp Asp Lys Ala Ser Asp Leu Leu Ser Gln
1 5 10 15
atc gtc agg aat gct gca tcc aat gac aaa ggg tct gac ctg atg act 96
Ile Val Arg Asn Ala Ala Ser Asn Asp Lys Gly Ser Asp Leu Met Thr
20 25 30
ctt gcc ctc agg gga tgc tgt aaa aat cct tac tgt ggt gcg tcg aaa 144
Leu Ala Leu Arg Gly Cys Cys Lys Asn Pro Tyr Cys Gly Ala Ser Lys
35 40 45
aca tat tgt ggt aga aga cgc tgatgctcca ggaccctctg aaccacgacg t 196
Thr Tyr Cys Gly Arg Arg Arg
50 55

<210> 272
<211> 55
<212> PRT
<213> Conus musicus

<400> 272
Ile Lys Asn Ala Ala Ala Asp Asp Lys Ala Ser Asp Leu Leu Ser Gln
1 5 10 15
Ile Val Arg Asn Ala Ala Ser Asn Asp Lys Gly Ser Asp Leu Met Thr
20 25 30
Leu Ala Leu Arg Gly Cys Cys Lys Asn Pro Tyr Cys Gly Ala Ser Lys
35 40 45
Thr Tyr Cys Gly Arg Arg Arg
50 55

<400> 276

Phe Asp Gly Arg Asn Ala Ala Ala Asp Tyr Lys Gly Ser Glu Leu Leu
 1 5 10 15

Ala Met Thr Val Arg Gly Gly Cys Cys Ser Tyr Pro Pro Cys Ile Ala
 20 25 30

Asn Asn Pro Leu Cys Ala Gly Arg Arg
 35 40

<210> 277

<211> 126

<212> DNA

<213> Conus radiatus

<220>

<221> CDS

<222> (1)..(123)

<400> 277

ttt gat ggc agg aat gcc gca gcc gac tac aaa ggg tct gaa ttg ctc 48
 Phe Asp Gly Arg Asn Ala Ala Ala Asp Tyr Lys Gly Ser Glu Leu Leu
 1 5 10 15

gct atg acc gtc agg gga gga tgc tgt tcc tat cct ccc tgt atc gca 96
 Ala Met Thr Val Arg Gly Gly Cys Cys Ser Tyr Pro Pro Cys Ile Ala
 20 25 30

aat aat cct ttt tgt gct gga aga cgc tga 126
 Asn Asn Pro Phe Cys Ala Gly Arg Arg
 35 40

<210> 278

<211> 41

<212> PRT

<213> Conus radiatus

<400> 278

Phe Asp Gly Arg Asn Ala Ala Ala Asp Tyr Lys Gly Ser Glu Leu Leu
 1 5 10 15

Ala Met Thr Val Arg Gly Gly Cys Cys Ser Tyr Pro Pro Cys Ile Ala
 20 25 30

Asn Asn Pro Phe Cys Ala Gly Arg Arg
 35 40

<210> 279

<211> 155

<212> DNA

<213> Conus virgo

<220>

<221> CDS

<222> (1)..(114)

<400> 279

tct tat gac agg tat gcc tcg ccc gtc gac aga gcg tct gcc ctg atc 48
 Ser Tyr Asp Arg Tyr Ala Ser Pro Val Asp Arg Ala Ser Ala Leu Ile
 1 5 10 15

gct cag gcc atc ctt cga gat tgc tgt tcc aat cct ccc tgt tcc caa 96
 Ala Gln Ala Ile Leu Arg Asp Cys Cys Ser Asn Pro Pro Cys Ser Gln
 20 25 30

aat aat cca gac tgt atg taaagacgct gcttgctcca ggaccctctg 144
 Asn Asn Pro Asp Cys Met
 35

aaccacgacg t 155

<210> 280
 <211> 38
 <212> PRT
 <213> Conus virgo

<400> 280
 Ser Tyr Asp Arg Tyr Ala Ser Pro Val Asp Arg Ala Ser Ala Leu Ile
 1 5 10 15

Ala Gln Ala Ile Leu Arg Asp Cys Cys Ser Asn Pro Pro Cys Ser Gln
 20 25 30

Asn Asn Pro Asp Cys Met
 35

<210> 281
 <211> 155
 <212> DNA
 <213> Conus virgo

<220>
 <221> CDS
 <222> (1)..(114)

<400> 281
 tct tat ggc agg tat gcc tca ccc gtc gac aga gcg tct gcc ctg atc 48
 Ser Tyr Gly Arg Tyr Ala Ser Pro Val Asp Arg Ala Ser Ala Leu Ile
 1 5 10 15

gct cag gcc atc ctt cga gat tgc tgc tcc aat cct cct tgt gcc cat 96
 Ala Gln Ala Ile Leu Arg Asp Cys Cys Ser Asn Pro Pro Cys Ala His
 20 25 30

aat aat cca gac tgt cgt taaagacgct gcttgctcca ggaccctctg 144
 Asn Asn Pro Asp Cys Arg
 35

aaccacgacg t 155

<210> 282
 <211> 38
 <212> PRT
 <213> Conus virgo

<400> 282
 Ser Tyr Gly Arg Tyr Ala Ser Pro Val Asp Arg Ala Ser Ala Leu Ile
 1 5 10 15

Ala Gln Ala Ile Leu Arg Asp Cys Cys Ser Asn Pro Pro Cys Ala His
 20 25 30

Asn Asn Pro Asp Cys Arg
35

<210> 283
<211> 126
<212> DNA
<213> Conus achatinus

<220>
<221> CDS
<222> (1)..(123)

<400> 283
tct gat ggc agg aat gcc gca gcc aac gac aaa gcg tct ggc atg agc 48
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Gly Met Ser
1 5 10 15
gcg ctg gcc gtc aat gaa tgc tgt acc aac cct gtc tgt cac gcg gaa 96
Ala Leu Ala Val Asn Glu Cys Cys Thr Asn Pro Val Cys His Ala Glu
20 25 30
cat caa gaa ctt tgt gct aga aga cgc tga 126
His Gln Glu Leu Cys Ala Arg Arg Arg
35 40

<210> 284
<211> 41
<212> PRT
<213> Conus achatinus

<400> 284
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Gly Met Ser
1 5 10 15
Ala Leu Ala Val Asn Glu Cys Cys Thr Asn Pro Val Cys His Ala Glu
20 25 30
His Gln Glu Leu Cys Ala Arg Arg Arg
35 40

<210> 285
<211> 126
<212> DNA
<213> Conus achatinus

<220>
<221> CDS
<222> (1)..(123)

<400> 285
tct gat ggc agg aat gcc gca gcc aac gac aaa gcg tct gac gtg atc 48
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Asp Val Ile
1 5 10 15
acg ctg gcc ctc aag gga tgc tgt tcc aac cct gtc tgt cac ttg gag 96
Thr Leu Ala Leu Lys Gly Cys Cys Ser Asn Pro Val Cys His Leu Glu
20 25 30
cat tca aac ctt tgt ggt aga aga cgc tga 126
His Ser Asn Leu Cys Gly Arg Arg Arg
35 40

<210> 286
 <211> 41
 <212> PRT
 <213> *Conus achatinus*

<400> 286
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Asp Val Ile
 1 5 10 15
 Thr Leu Ala Leu Lys Gly Cys Cys Ser Asn Pro Val Cys His Leu Glu
 20 25 30
 His Ser Asn Leu Cys Gly Arg Arg Arg
 35 40

<210> 287
 <211> 126
 <212> DNA
 <213> *Conus achatinus*

<220>
 <221> CDS
 <222> (1)..(123)

<400> 287
 tct gat ggc agg aat gcc gca gcc aac gac aaa gcg tct ggc atg agc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Gly Met Ser
 1 5 10 15
 gcg ctg gcc gtc aat gaa tgc tgt acc aac cct gtc tgt cac gtg gaa 96
 Ala Leu Ala Val Asn Glu Cys Cys Thr Asn Pro Val Cys His Val Glu
 20 25 30
 cat caa gaa ctt tgt gct aga aga cgc tga 126
 His Gln Glu Leu Cys Ala Arg Arg Arg
 35 40

<210> 288
 <211> 41
 <212> PRT
 <213> *Conus achatinus*

<400> 288
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Gly Met Ser
 1 5 10 15
 Ala Leu Ala Val Asn Glu Cys Cys Thr Asn Pro Val Cys His Val Glu
 20 25 30
 His Gln Glu Leu Cys Ala Arg Arg Arg
 35 40

<210> 289
 <211> 220
 <212> DNA
 <213> *Conus ammiralis*

<220>
 <221> CDS
 <222> (1)..(180)

<400> 289

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ttc act tca gat cgt gca ttt cgt ggc agg aat gcc gca gcc aaa gcg 96
 Phe Thr Ser Asp Arg Ala Phe Arg Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30

tct ggc ctg gtc ggt ctg acc gac aag agg caa gaa tgc tgt tct tat 144
 Ser Gly Leu Val Gly Leu Thr Asp Lys Arg Gln Glu Cys Cys Ser Tyr
 35 40 45

cct gcc tgt aac cta gat cat cca gaa ctt tgt ggt tgaagacgct 190
 Pro Ala Cys Asn Leu Asp His Pro Glu Leu Cys Gly
 50 55 60

gatgctccag gaccctctga accacgacgt 220

<210> 290

<211> 60

<212> PRT

<213> Conus ammiralis

<400> 290

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Phe Thr Ser Asp Arg Ala Phe Arg Gly Arg Asn Ala Ala Ala Lys Ala
 20 25 30

Ser Gly Leu Val Gly Leu Thr Asp Lys Arg Gln Glu Cys Cys Ser Tyr
 35 40 45

Pro Ala Cys Asn Leu Asp His Pro Glu Leu Cys Gly
 50 55 60

<210> 291

<211> 223

<212> DNA

<213> Conus ammiralis

<220>

<221> CDS

<222> (1)..(192)

<400> 291

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc act gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

tcc act tca ggt cgt cgt gca ttt cgt ggc agg aat gcc gca gcc aaa 96
 Ser Thr Ser Gly Arg Arg Ala Phe Arg Gly Arg Asn Ala Ala Ala Lys
 20 25 30

gcg tct gga ctg gtc ggt ctg act gac agg aga cca gaa tgc tgt agt 144
 Ala Ser Gly Leu Val Gly Leu Thr Asp Arg Arg Pro Glu Cys Cys Ser
 35 40 45

gat cct cgc tgt aac tcg act cat cca gaa ctt tgt ggt gga aga cgc 192
 Asp Pro Arg Cys Asn Ser Thr His Pro Glu Leu Cys Gly Gly Arg Arg
 50 55 60

tgatgctcca ggaccctctg aaccacgacg t

223

<210> 292
 <211> 64
 <212> PRT
 <213> Conus ammiralis

<400> 292
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15
 Ser Thr Ser Gly Arg Arg Ala Phe Arg Gly Arg Asn Ala Ala Ala Lys
 20 25 30
 Ala Ser Gly Leu Val Gly Leu Thr Asp Arg Arg Pro Glu Cys Cys Ser
 35 40 45
 Asp Pro Arg Cys Asn Ser Thr His Pro Glu Leu Cys Gly Gly Arg Arg
 50 55 60

<210> 293
 <211> 151
 <212> DNA
 <213> Conus arenatus

<220>
 <221> CDS
 <222> (1)...(120)

<400> 293
 tct gat ggc agg aat gcc gca gcc aac gcg ttt gac ctg atc gat ctg 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala Phe Asp Leu Ile Asp Leu
 1 5 10 15
 acc gcc agg cta aat tgc tgt atg att ccc ccc tgt tgg aag aaa tat 96
 Thr Ala Arg Leu Asn Cys Cys Met Ile Pro Pro Cys Trp Lys Lys Tyr
 20 25 30
 gga gac aga tgt agt gaa gta cgc tgatgctcca ggaccctctg aaccacgacg 150
 Gly Asp Arg Cys Ser Glu Val Arg
 35 40
 t 151

<210> 294
 <211> 40
 <212> PRT
 <213> Conus arenatus

<400> 294
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala Phe Asp Leu Ile Asp Leu
 1 5 10 15
 Thr Ala Arg Leu Asn Cys Cys Met Ile Pro Pro Cys Trp Lys Lys Tyr
 20 25 30
 Gly Asp Arg Cys Ser Glu Val Arg
 35 40

<210> 295

<211> 126
 <212> DNA
 <213> Conus arenatus

<220>
 <221> CDS
 <222> (1)..(93)

<400> 295
 tct gat ggc agg aat gcc gca cgc aaa gcg ttt ggc tgc tgc gac tta 48
 Ser Asp Gly Arg Asn Ala Ala Arg Lys Ala Phe Gly Cys Cys Asp Leu
 1 5 10 15
 ata ccc tgt ttg gag aga tat ggt aac aga tgt aat gaa gtg cac 93
 Ile Pro Cys Leu Glu Arg Tyr Gly Asn Arg Cys Asn Glu Val His
 20 25 30
 tgatgctcca ggaccctctg aaccacgcga cgt 126

<210> 296
 <211> 31
 <212> PRT
 <213> Conus arenatus

<400> 296
 Ser Asp Gly Arg Asn Ala Ala Arg Lys Ala Phe Gly Cys Cys Asp Leu
 1 5 10 15
 Ile Pro Cys Leu Glu Arg Tyr Gly Asn Arg Cys Asn Glu Val His
 20 25 30

<210> 297
 <211> 151
 <212> DNA
 <213> Conus arenatus

<220>
 <221> CDS
 <222> (1)..(120)

<400> 297
 tct gat ggc agc aat gcc gca gcc aac gag ttt gac ctg atc gct ctg 48
 Ser Asp Gly Ser Asn Ala Ala Ala Asn Glu Phe Asp Leu Ile Ala Leu
 1 5 10 15
 acc gcc agg cta ggt tgc tgt aac gtt aca ccc tgt tgg gag aaa tat 96
 Thr Ala Arg Leu Gly Cys Cys Asn Val Thr Pro Cys Trp Glu Lys Tyr
 20 25 30
 gga gac aaa tgt aat gaa gta cgc tgatgcttca ggaccctctg aaccacgcag 150
 Gly Asp Lys Cys Asn Glu Val Arg
 35 40
 t 151

<210> 298
 <211> 40
 <212> PRT
 <213> Conus arenatus

<400> 298

Ser Asp Gly Ser Asn Ala Ala Ala Asn Glu Phe Asp Leu Ile Ala Leu
 1 5 10 15

Thr Ala Arg Leu Gly Cys Cys Asn Val Thr Pro Cys Trp Glu Lys Tyr
 20 25 30

Gly Asp Lys Cys Asn Glu Val Arg
 35 40

<210> 299

<211> 148

<212> DNA

<213> Conus arenatus

<220>

<221> CDS

<222> (1)..(117)

<400> 299

tct gat ggc agg aat gtc gca gca aaa gcg ttt cac cgg atc ggc cgg 48
 Ser Asp Gly Arg Asn Val Ala Ala Lys Ala Phe His Arg Ile Gly Arg
 1 5 10 15

acc atc agg gat gaa tgc tgt tcc aat cct gcc tgt agg gtg aat aat 96
 Thr Ile Arg Asp Glu Cys Cys Ser Asn Pro Ala Cys Arg Val Asn Asn
 20 25 30

cca cac gtt tgt aga cga cgc tgatgctcca ggaccctctg aaccacgacg t 148
 Pro His Val Cys Arg Arg Arg
 35

<210> 300

<211> 39

<212> PRT

<213> Conus arenatus

<400> 300

Ser Asp Gly Arg Asn Val Ala Ala Lys Ala Phe His Arg Ile Gly Arg
 1 5 10 15

Thr Ile Arg Asp Glu Cys Cys Ser Asn Pro Ala Cys Arg Val Asn Asn
 20 25 30

Pro His Val Cys Arg Arg Arg
 35

<210> 301

<211> 151

<212> DNA

<213> Conus arenatus

<220>

<221> CDS

<222> (1)..(120)

<400> 301

tct gat ggc agg aat gcc gca gcc aac gcg ttt gac ctg atg cct ctg 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala Phe Asp Leu Met Pro Leu
 1 5 10 15

acc gcc agg cta aat tgc tgt agc att ccc gcc tgt tgg aac gaa tat 96

Thr Ala Arg Leu Asn Cys Cys Ser Ile Pro Gly Cys Trp Asn Glu Tyr
 20 25 30

aaa gac aga tgt agt aaa gta cgc tgatgctcca ggaccctctg aaccacgacg 150
 Lys Asp Arg Cys Ser Lys Val Arg
 35 40

t 151

<210> 302
 <211> 40
 <212> PRT
 <213> Conus arenatus

<400> 302
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala Phe Asp Leu Met Pro Leu
 1 5 10 15

Thr Ala Arg Leu Asn Cys Cys Ser Ile Pro Gly Cys Trp Asn Glu Tyr
 20 25 30

Lys Asp Arg Cys Ser Lys Val Arg
 35 40

<210> 303
 <211> 157
 <212> DNA
 <213> Conus aurisiacus

<220>
 <221> CDS
 <222> (52)..(126)

<400> 303
 tctgatggca ggaatgccgc agccgacgac aaagcgtctg acctggtcgc t ctg gtc 57
 Leu Val
 1

gtc agg gga gga tgc tgt tcc cac cct gtc tgt tac ttt aat aat cca 105
 Val Arg Gly Gly Cys Cys Ser His Pro Val Cys Tyr Phe Asn Asn Pro
 5 10 15

caa atg tgt cgt gga aga cgc tgatgctcca ggaccctctg aaccacgacg t 157
 Gln Met Cys Arg Gly Arg Arg
 20 25

<210> 304
 <211> 25
 <212> PRT
 <213> Conus aurisiacus

<400> 304
 Leu Val Val Arg Gly Gly Cys Cys Ser His Pro Val Cys Tyr Phe Asn
 1 5 10 15

Asn Pro Gln Met Cys Arg Gly Arg Arg
 20 25

<210> 305
 <211> 157

<212> DNA
 <213> Conus aurisiacus

<220>
 <221> CDS
 <222> (52)..(126)

<400> 305
 tctgatggca ggaatgccgc agccgacgac aaagcgtctg acctggctgc t ctg gcc 57
 Leu Ala
 1
 gtc agg gga gga tgc tgt tcc cac cct gtc tgt aac ttg aat aat cca 105
 Val Arg Gly Gly Cys Cys Ser His Pro Val Cys Asn Leu Asn Asn Pro
 5 10 15
 caa atg tgt cgt gga aga cgc tgatgctcca ggaccctctg aaccacgacg t 157
 Gln Met Cys Arg Gly Arg Arg
 20 25

<210> 306
 <211> 25
 <212> PRT
 <213> Conus aurisiacus

<400> 306
 Leu Ala Val Arg Gly Gly Cys Cys Ser His Pro Val Cys Asn Leu Asn
 1 5 10 15
 Asn Pro Gln Met Cys Arg Gly Arg Arg
 20 25

<210> 307
 <211> 157
 <212> DNA
 <213> Conus betulinus

<220>
 <221> CDS
 <222> (1)..(117)

<400> 307
 ttt cgt ggc agg aat ccc gca gcc aac gac aaa agg tct gac ctg gcc 48
 Phe Arg Gly Arg Asn Pro Ala Ala Asn Asp Lys Arg Ser Asp Leu Ala
 1 5 10 15
 gct ctg agc gtc agg gga gga tgc tgt tcc cat cct gcc tgt agc gtg 96
 Ala Leu Ser Val Arg Gly Gly Cys Cys Ser His Pro Ala Cys Ser Val
 20 25 30
 act cat cca gag ctt tgt ggc tgaagacgct gatgccccag gaccctctga 147
 Thr His Pro Glu Leu Cys Gly
 35
 accacgacgt 157

<210> 308
 <211> 39
 <212> PRT
 <213> Conus betulinus

<400> 308

Phe Arg Gly Arg Asn Pro Ala Ala Asn Asp Lys Arg Ser Asp Leu Ala
 1 5 10 15

Ala Leu Ser Val Arg Gly Gly Cys Cys Ser His Pro Ala Cys Ser Val
 20 25 30

Thr His Pro Glu Leu Cys Gly
 35

<210> 309

<211> 151

<212> DNA

<213> Conus betulinus

<220>

<221> CDS

<222> (1)..(120)

<400> 309

tct gat ggc ggg aat gcc gca gcc aaa gcg tct gac ctg atc gct cag 48
 Ser Asp Gly Gly Asn Ala Ala Ala Lys Ala Ser Asp Leu Ile Ala Gln
 1 5 10 15

acc atc agg gga gga tgc tgt tcc tat cct gcc tgt agc gtg gaa cat 96
 Thr Ile Arg Gly Gly Cys Cys Ser Tyr Pro Ala Cys Ser Val Glu His
 20 25 30

caa gac ott tgt gat gga aga cgc tgatgctcca ggaccctctg aaccacgacg 150
 Gln Asp Leu Cys Asp Gly Arg Arg
 35 40

t 151

<210> 310

<211> 40

<212> PRT

<213> Conus betulinus

<400> 310

Ser Asp Gly Gly Asn Ala Ala Ala Lys Ala Ser Asp Leu Ile Ala Gln
 1 5 10 15

Thr Ile Arg Gly Gly Cys Cys Ser Tyr Pro Ala Cys Ser Val Glu His
 20 25 30

Gln Asp Leu Cys Asp Gly Arg Arg
 35 40

<210> 311

<211> 114

<212> DNA

<213> Conus characteristicus

<220>

<221> CDS

<222> (1)..(111)

<400> 311

tct tat ggc agg aat gcc gca gcc aaa gcg ttt gaa gtg agt tgc tgt 48
 Ser Tyr Gly Arg Asn Ala Ala Ala Lys Ala Phe Glu Val Ser Cys Cys

1 5 10 15

gtc gtt cgc ccc tgt tgg att cgc tat caa gag gaa tgt ctt gaa gca 96
 Val Val Arg Pro Cys Trp Ile Arg Tyr Gln Glu Glu Cys Leu Glu Ala
 20 25 30

gat ccc agg acc ctc tga 114
 Asp Pro Arg Thr Leu
 35

<210> 312
 <211> 37
 <212> PRT
 <213> Conus characteristicus

<400> 312
 Ser Tyr Gly Arg Asn Ala Ala Ala Lys Ala Phe Glu Val Ser Cys Cys
 1 5 10 15

Val Val Arg Pro Cys Trp Ile Arg Tyr Gln Glu Glu Cys Leu Glu Ala
 20 25 30

Asp Pro Arg Thr Leu
 35

<210> 313
 <211> 123
 <212> DNA
 <213> Conus characteristicus

<220>
 <221> CDS
 <222> (1)..(120)

<400> 313
 tct gat ggc agg aat gcc gca gcc aac gcc ctt gac ctg atc act ctg 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala Leu Asp Leu Ile Thr Leu
 1 5 10 15

atc gcc agg caa aat tgc tgt agc att ccc ggc tgt tgg gag aaa tat 96
 Ile Ala Arg Gln Asn Cys Cys Ser Ile Pro Gly Cys Trp Glu Lys Tyr
 20 25 30

gga gac aaa tgt agt gaa gta cgc tga 123
 Gly Asp Lys Cys Ser Glu Val Arg
 35 40

<210> 314
 <211> 40
 <212> PRT
 <213> Conus characteristicus

<400> 314
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Ala Leu Asp Leu Ile Thr Leu
 1 5 10 15

Ile Ala Arg Gln Asn Cys Cys Ser Ile Pro Gly Cys Trp Glu Lys Tyr
 20 25 30

Gly Asp Lys Cys Ser Glu Val Arg
 35 40

<210> 315
 <211> 154
 <212> DNA
 <213> Conus catus

<220>
 <221> CDS
 <222> (1)..(123)

<400> 315
 tct gat ggc agg aat gaa gca gcc aac gac gaa gcg tct gac gtg atc 48
 Ser Asp Gly Arg Asn Glu Ala Ala Asn Asp Glu Ala Ser Asp Val Ile
 1 5 10 15
 gag ctg gcc ctc aag gga tgc tgt tcc aac cct gtc tgt cac ttg gag 96
 Glu Leu Ala Leu Lys Gly Cys Cys Ser Asn Pro Val Cys His Leu Glu
 20 25 30
 cat cca aac gct tgt ggt aga aga cgc tgatgctcca ggaccctctg 143
 His Pro Asn Ala Cys Gly Arg Arg Arg
 35 40
 aaccacgacg t 154

<210> 316
 <211> 41
 <212> PRT
 <213> Conus catus

<400> 316
 Ser Asp Gly Arg Asn Glu Ala Ala Asn Asp Glu Ala Ser Asp Val Ile
 1 5 10 15
 Glu Leu Ala Leu Lys Gly Cys Cys Ser Asn Pro Val Cys His Leu Glu
 20 25 30
 His Pro Asn Ala Cys Gly Arg Arg Arg
 35 40

<210> 317
 <211> 154
 <212> DNA
 <213> Conus catus

<220>
 <221> CDS
 <222> (1)..(123)

<400> 317
 tct gat ggc agg aat gcc gca gcc aac gac aaa gcg tct gac ctg gtc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Asp Leu Val
 1 5 10 15
 gct ctg gcc gtc agg gga tgc tgt tcc aac cct atc tgt tac ttt aat 96
 Ala Leu Ala Val Arg Gly Cys Cys Ser Asn Pro Ile Cys Tyr Phe Asn
 20 25 30
 aat cca cga att tgt cgt gga aga cgc tgatgctcca ggaccctctg 143
 Asn Pro Arg Ile Cys Arg Gly Arg Arg
 35 40

aaccacgacg t

154

<210> 318
 <211> 41
 <212> PRT
 <213> Conus catus

<400> 318
 Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Asp Leu Val
 1 5 10 15
 Ala Leu Ala Val Arg Gly Cys Cys Ser Asn Pro Ile Cys Tyr Phe Asn
 20 25 30
 Asn Pro Arg Ile Cys Arg Gly Arg Arg
 35 40

<210> 319
 <211> 111
 <212> DNA
 <213> Conus episcopatus

<220>
 <221> CDS
 <222> (1)..(108)

<400> 319
 tct cat ggc agg aat gcc gca cgc aaa gcg tct gac ctg atc gct ctg 48
 Ser His Gly Arg Asn Ala Ala Arg Lys Ala Ser Asp Leu Ile Ala Leu
 1 5 10 15
 acc gtc agg gaa tgc tgt tct cag cct ccc tgt cgc tgg aaa cat cca 96
 Thr Val Arg Glu Cys Cys Ser Gln Pro Pro Cys Arg Trp Lys His Pro
 20 25 30
 gaa ctt tgt agt tga 111
 Glu Leu Cys Ser
 35

<210> 320
 <211> 36
 <212> PRT
 <213> Conus episcopatus

<400> 320
 Ser His Gly Arg Asn Ala Ala Arg Lys Ala Ser Asp Leu Ile Ala Leu
 1 5 10 15
 Thr Val Arg Glu Cys Cys Ser Gln Pro Pro Cys Arg Trp Lys His Pro
 20 25 30
 Glu Leu Cys Ser
 35

<210> 321
 <211> 151
 <212> DNA
 <213> Conus geographus

<220>

<221> CDS
 <222> (1)..(120)

<400> 321

tct gat ggc agg aat gac gca gcc aaa gcg ttt gac ctg ata tct tcg	48
Ser Asp Gly Arg Asn Asp Ala Ala Lys Ala Phe Asp Leu Ile Ser Ser	
1 5 10 15	
acc gtc aag aaa gga tgc tgt tcc cat cct gcc tgt gcg ggg aat aat	96
Thr Val Lys Lys Gly Cys Cys Ser His Pro Ala Cys Ala Gly Asn Asn	
20 25 30	
caa cat att tgt ggc cga aga cgc tgatgctcca ggaccctctg aaccacgacg	150
Gln His Ile Cys Gly Arg Arg Arg	
35 40	

t 151

<210> 322
 <211> 40
 <212> PRT
 <213> Conus geographus

<400> 322

Ser Asp Gly Arg Asn Asp Ala Ala Lys Ala Phe Asp Leu Ile Ser Ser	
1 5 10 15	
Thr Val Lys Lys Gly Cys Cys Ser His Pro Ala Cys Ala Gly Asn Asn	
20 25 30	
Gln His Ile Cys Gly Arg Arg Arg	
35 40	

<210> 323
 <211> 154
 <212> DNA
 <213> Conus geographus

<220>
 <221> CDS
 <222> (1)..(123)

<400> 323

tct gat ggc agg aat gcc gca gcc aac gac caa gcg tct gac ctg atg	48
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Gln Ala Ser Asp Leu Met	
1 5 10 15	
gct gcg acc gtc agg gga tgc tgt gcc gtt cct tcc tgt cgc ctc cgt	96
Ala Ala Thr Val Arg Gly Cys Cys Ala Val Pro Ser Cys Arg Leu Arg	
20 25 30	
aat cca gac ctt tgt ggt gga gga cgc tgatgctcca ggaccctctg	143
Asn Pro Asp Leu Cys Gly Gly Gly Arg	
35 40	

aaccacgacg t 154

<210> 324
 <211> 41
 <212> PRT
 <213> Conus geographus

<400> 324

Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Gln Ala Ser Asp Leu Met
 1 5 10 15

Ala Ala Thr Val Arg Gly Cys Cys Ala Val Pro Ser Cys Arg Leu Arg
 20 25 30

Asn Pro Asp Leu Cys Gly Gly Gly Arg
 35 40

<210> 325

<211> 120

<212> DNA

<213> *Conus imperialis*

<220>

<221> CDS

<222> (1)..(117)

<400> 325

ctt gat gaa agg aat gcc gca gcc gac gac aaa gcg tct gac ctg atc 48
 Leu Asp Glu Arg Asn Ala Ala Ala Asp Asp Lys Ala Ser Asp Leu Ile
 1 5 10 15

gct caa atc gtc agg aga gga tgc tgt tcc cat cct gcc tgt aac gtg 96
 Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Ala Cys Asn Val
 20 25 30

aat aat cca cac att tgt ggt tga 120
 Asn Asn Pro His Ile Cys Gly
 35

<210> 326

<211> 39

<212> PRT

<213> *Conus imperialis*

<400> 326

Leu Asp Glu Arg Asn Ala Ala Ala Asp Asp Lys Ala Ser Asp Leu Ile
 1 5 10 15

Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Ala Cys Asn Val
 20 25 30

Asn Asn Pro His Ile Cys Gly
 35

<210> 327

<211> 142

<212> DNA

<213> *Conus lividus*

<220>

<221> CDS

<222> (1)..(111)

<400> 327

tct gat ggc agg aat act gca gcc aaa gtc aaa tat tct aag acg ccg 48
 Ser Asp Gly Arg Asn Thr Ala Ala Lys Val Lys Tyr Ser Lys Thr Pro
 1 5 10 15

gag gaa tgc tgt ccc aat cct ccc tgt ttc gcg aca aat tcg gat att 96
 Glu Glu Cys Cys Pro Asn Pro Pro Cys Phe Ala Thr Asn Ser Asp Ile
 20 25 30

tgt ggc gga aga cgc tgatgctcca ggaccctctg aaccacgacg t 142
 Cys Gly Gly Arg Arg
 35

<210> 328
 <211> 37
 <212> PRT
 <213> Conus lividus

<400> 328
 Ser Asp Gly Arg Asn Thr Ala Ala Lys Val Lys Tyr Ser Lys Thr Pro
 1 5 10 15
 Glu Glu Cys Cys Pro Asn Pro Pro Cys Phe Ala Thr Asn Ser Asp Ile
 20 25 30
 Cys Gly Gly Arg Arg
 35

<210> 329
 <211> 157
 <212> DNA
 <213> Conus lividus

<220>
 <221> CDS
 <222> (1)..(117)

<400> 329
 tct aat ggc agg aat gcc gca gcc aaa ttc aaa gcg cct gcc ctg atg 48
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 aag cgg acc gtc agg gat gct tgc tgt tca gac cct cgc tgt tcc ggg 96
 Lys Arg Thr Val Arg Asp Ala Cys Cys Ser Asp Pro Arg Cys Ser Gly
 20 25 30
 aaa cat caa gac ctg tgt ggc tgaagacgct gatgctccag gaccctctga 147
 Lys His Gln Asp Leu Cys Gly
 35
 accacgacgt 157

<210> 330
 <211> 39
 <212> PRT
 <213> Conus lividus

<400> 330
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 Lys Arg Thr Val Arg Asp Ala Cys Cys Ser Asp Pro Arg Cys Ser Gly
 20 25 30
 Lys His Gln Asp Leu Cys Gly
 35

<210> 331
 <211> 157
 <212> DNA
 <213> Conus lividus

<220>
 <221> CDS
 <222> (1)..(117)

<400> 331
 tct aat ggc agg aat gcc gca gcc aaa ttc aaa gcg cct gcc ctg atg 48
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 gag ctg acc gtc agg gaa gat tgc tgt tca gac cct cgc tgt tcc gtg 96
 Glu Leu Thr Val Arg Glu Asp Cys Cys Ser Asp Pro Arg Cys Ser Val
 20 25 30
 gga cat caa gac ctg tgt ggc tgaagacgct gatgctccag gaccctctga 147
 Gly His Gln Asp Leu Cys Gly
 35
 accacgacgt. 157

<210> 332
 <211> 39
 <212> PRT
 <213> Conus lividus

<400> 332
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 Glu Leu Thr Val Arg Glu Asp Cys Cys Ser Asp Pro Arg Cys Ser Val
 20 25 30
 Gly His Gln Asp Leu Cys Gly
 35

<210> 333
 <211> 157
 <212> DNA
 <213> Conus lividus

<220>
 <221> CDS
 <222> (1)..(126)

<400> 333
 gca ttt gat ggc agg aat gct gca gcc agc gac aaa gcg tcc gag ctg 48
 Ala Phe Asp Gly Arg Asn Ala Ala Ala Ser Asp Lys Ala Ser Glu Leu
 1 5 10 15
 atg gct ctg gcc gtc agg gga tgc tgt tcc cat cct gcc tgt gct ggg 96
 Met Ala Leu Ala Val Arg Gly Cys Cys Ser His Pro Ala Cys Ala Gly
 20 25 30
 agt aat gca cat atc tgt ggc aga aga cgc tgatgctcca ggaccctctg 146
 Ser Asn Ala His Ile Cys Gly Arg Arg
 35 40

aaccacgacg t

157

<210> 334
 <211> 42
 <212> PRT
 <213> Conus lividus

<400> 334
 Ala Phe Asp Gly Arg Asn Ala Ala Ala Ser Asp Lys Ala Ser Glu Leu
 1 5 10 15
 Met Ala Leu Ala Val Arg Gly Cys Cys Ser His Pro Ala Cys Ala Gly
 20 25 30
 Ser Asn Ala His Ile Cys Gly Arg Arg Arg
 35 40

<210> 335
 <211> 157
 <212> DNA
 <213> Conus lividus

<220>
 <221> CDS
 <222> (1)..(117)

<400> 335
 tct aat ggc agg aat gcc gca gcc aaa ttc aaa gcg cct gcc ctg atg 48
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 aag ctg acc gtc agg gag gat tgc tgt tca gac cct cgc tgt tcc gtg 96
 Lys Leu Thr Val Arg Glu Asp Cys Cys Ser Asp Pro Arg Cys Ser Val
 20 25 30
 gga cat caa gac atg tgt ggc tgaagacgct gatgctccag gaccctctga 147
 Gly His Gln Asp Met Cys Gly
 35

atcacgacgt

157

<210> 336
 <211> 39
 <212> PRT
 <213> Conus lividus

<400> 336
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 Lys Leu Thr Val Arg Glu Asp Cys Cys Ser Asp Pro Arg Cys Ser Val
 20 25 30
 Gly His Gln Asp Met Cys Gly
 35

<210> 337
 <211> 154
 <212> DNA
 <213> Conus lividus

<220>

<221> CDS

<222> (1)..(114)

<400> 337

ttt	gaa	tgc	agg	aat	gct	gca	ggc	aac	gac	aaa	gcg	act	gac	ctg	atg	48
Phe	Glu	Cys	Arg	Asn	Ala	Ala	Gly	Asn	Asp	Lys	Ala	Thr	Asp	Leu	Met	
1				5					10					15		

gct	ctg	act	gtc	agg	gga	tgc	tgt	tcc	cat	cct	gcc	tgt	gct	ggg	aat	96
Ala	Leu	Thr	Val	Arg	Gly	Cys	Cys	Ser	His	Pro	Ala	Cys	Ala	Gly	Asn	
			20					25						30		

aat	cca	cat	atc	tgc	ggc	tgaagacgct	gatgctccag	gaccctctga	144
Asn	Pro	His	Ile	Cys	Gly				
			35						

accacgacgt	154
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<210> 338

<211> 38

<212> PRT

<213> Conus lividus

<400> 338

Phe	Glu	Cys	Arg	Asn	Ala	Ala	Gly	Asn	Asp	Lys	Ala	Thr	Asp	Leu	Met
1				5					10					15	

Ala	Leu	Thr	Val	Arg	Gly	Cys	Cys	Ser	His	Pro	Ala	Cys	Ala	Gly	Asn
			20					25						30	

Asn	Pro	His	Ile	Cys	Gly
			35		

<210> 339

<211> 154

<212> DNA

<213> Conus lividus

<220>

<221> CDS

<222> (1)..(114)

<400> 339

ttt	gat	ggc	agg	aac	gcc	gca	gcc	aac	aac	aaa	gcg	act	gat	ctg	atg	48
Phe	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Asn	Asn	Lys	Ala	Thr	Asp	Leu	Met	
1				5					10					15		

gct	ctg	act	gtc	aga	gga	tgc	tgt	ggc	aat	cct	tca	tgt	agc	atc	cat	96
Ala	Leu	Thr	Val	Arg	Gly	Cys	Cys	Gly	Asn	Pro	Ser	Cys	Ser	Ile	His	
			20					25						30		

att	cct	tac	gtt	tgt	aat	tagagacact	gatgctccag	gaccctctga	144
Ile	Pro	Tyr	Val	Cys	Asn				
			35						

accacgacgt	154
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<210> 340

<211> 38

<212> PRT

<213> Conus lividus

<400> 340

Phe Asp Gly Arg Asn Ala Ala Ala Asn Asn Lys Ala Thr Asp Leu Met
 1 5 10 15
 Ala Leu Thr Val Arg Gly Cys Cys Gly Asn Pro Ser Cys Ser Ile His
 20 25 30
 Ile Pro Tyr Val Cys Asn
 35

<210> 341

<211> 157

<212> DNA

<213> Conus lividus

<220>

<221> CDS

<222> (1)..(126)

<400> 341

tct aat ggc agg aat gcc gca gcc aaa ttc aaa gcg cct gcc ctg atg 48
 Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 aag cgg acc gac agc gaa gaa tgc tgt tta gac tct cgc tgt gcc ggg 96
 Lys Arg Thr Asp Ser Glu Glu Cys Cys Leu Asp Ser Arg Cys Ala Gly
 20 25 30
 caa cat caa gac ctg tgt ggc gga aga cgc tgatgctcca ggaccctctg 146
 Gln His Gln Asp Leu Cys Gly Gly Arg Arg
 35 40
 aaccacgacg t 157

<210> 342

<211> 42

<212> PRT

<213> Conus lividus

<400> 342

Ser Asn Gly Arg Asn Ala Ala Ala Lys Phe Lys Ala Pro Ala Leu Met
 1 5 10 15
 Lys Arg Thr Asp Ser Glu Glu Cys Cys Leu Asp Ser Arg Cys Ala Gly
 20 25 30
 Gln His Gln Asp Leu Cys Gly Gly Arg Arg
 35 40

<210> 343

<211> 126

<212> DNA

<213> Conus marmoreus

<220>

<221> CDS

<222> (1)..(123)

<400> 343

tct gat ggc agg aat gcc gca gcc aag gac aaa gcg tct gac ctg gtc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Lys Asp Lys Ala Ser Asp Leu Val
 1 5 10 15

gct ctg acc gtc aag gga tgc tgt tct aat cct ccc tgt tac gcg aat 96
 Ala Leu Thr Val Lys Gly Cys Cys Ser Asn Pro Pro Cys Tyr Ala Asn
 20 25 30

aat caa gcc tat tgt aat gga aga cgc tga 126
 Asn Gln Ala Tyr Cys Asn Gly Arg Arg
 35 40

<210> 344
 <211> 41
 <212> PRT
 <213> Conus marmoreus

<400> 344
 Ser Asp Gly Arg Asn Ala Ala Ala Lys Asp Lys Ala Ser Asp Leu Val
 1 5 10 15
 Ala Leu Thr Val Lys Gly Cys Cys Ser Asn Pro Pro Cys Tyr Ala Asn
 20 25 30
 Asn Gln Ala Tyr Cys Asn Gly Arg Arg
 35 40

<210> 345
 <211> 117
 <212> DNA
 <213> Conus marmoreus

<220>
 <221> CDS
 <222> (1)..(114)

<400> 345
 tct gat ggc agg aat gcc gca gcc aag gac aaa gcg tct gac ctg gtc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Lys Asp Lys Ala Ser Asp Leu Val
 1 5 10 15
 gct ctg acc gtc aag gga tgc tgt tct cat cct gcc tgt agc gtg aat 96
 Ala Leu Thr Val Lys Gly Cys Cys Ser His Pro Ala Cys Ser Val Asn
 20 25 30
 aat cca gac att tgt ggt tga 117
 Asn Pro Asp Ile Cys Gly
 35

<210> 346
 <211> 38
 <212> PRT
 <213> Conus marmoreus

<400> 346
 Ser Asp Gly Arg Asn Ala Ala Ala Lys Asp Lys Ala Ser Asp Leu Val
 1 5 10 15
 Ala Leu Thr Val Lys Gly Cys Cys Ser His Pro Ala Cys Ser Val Asn
 20 25 30

Asn Pro Asp Ile Cys Gly
35

<210> 347
<211> 145
<212> DNA
<213> Conus musicus

<220>
<221> CDS
<222> (1)..(114)

<400> 347
tct gat ggc agg aat gct gca gcc aac aac aaa gtg gct ttg acc atg 48
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asn Lys Val Ala Leu Thr Met
1 5 10 15

agg gga aaa tgc tgt atc aat gat gcg tgt cgc tcg aaa cat cca cag 96
Arg Gly Lys Cys Cys Ile Asn Asp Ala Cys Arg Ser Lys His Pro Gln
20 25 30

tac tgt tct gga aga cgc tgatactcca ggaccctctg aaccacgacg t 145
Tyr Cys Ser Gly Arg Arg
35

<210> 348
<211> 38
<212> PRT
<213> Conus musicus

<400> 348
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asn Lys Val Ala Leu Thr Met
1 5 10 15

Arg Gly Lys Cys Cys Ile Asn Asp Ala Cys Arg Ser Lys His Pro Gln
20 25 30

Tyr Cys Ser Gly Arg Arg
35

<210> 349
<211> 154
<212> DNA
<213> Conus musicus

<220>
<221> CDS
<222> (1)..(123)

<400> 349
tct gat ggc agg aat gct gca gcc aac gac aaa gtg tct gac cag atg 48
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Val Ser Asp Gln Met
1 5 10 15

gct ctg gtt gtc agg gga tgc tgt tac aat att gcc tgt aga att aat 96
Ala Leu Val Val Arg Gly Cys Cys Tyr Asn Ile Ala Cys Arg Ile Asn
20 25 30

aat cca cgg tac tgt cgt gga aaa cgc tgatgttcca ggaccctctg 143
Asn Pro Arg Tyr Cys Arg Gly Lys Arg
35 40

aaccacgacg t

154

<210> 350

<211> 41

<212> PRT

<213> Conus musicus

<400> 350

Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Val Ser Asp Gln Met
 1 5 10 15

Ala Leu Val Val Arg Gly Cys Cys Tyr Asn Ile Ala Cys Arg Ile Asn
 20 25 30

Asn Pro Arg Tyr Cys Arg Gly Lys Arg
 35 40

<210> 351

<211> 154

<212> DNA

<213> Conus obscurus

<220>

<221> CDS

<222> (52)..(123)

<400> 351

tctgaaggca ggaatgccgc agccaacgac aaagcgtctg acctgatggc t ctg aac 57
 Leu Asn
 1

gtc agg gga tgc tgt tcc cat cct gtc tgt cgc ttc aat tat cca aaa 105
 Val Arg Gly Cys Cys Ser His Pro Val Cys Arg Phe Asn Tyr Pro Lys
 5 10 15

tat tgt ggt gga aga cgc tgatggtcca ggacctctg aaccacgacg t 154
 Tyr Cys Gly Gly Arg Arg
 20

<210> 352

<211> 24

<212> PRT

<213> Conus obscurus

<400> 352

Leu Asn Val Arg Gly Cys Cys Ser His Pro Val Cys Arg Phe Asn Tyr
 1 5 10 15

Pro Lys Tyr Cys Gly Gly Arg Arg
 20

<210> 353

<211> 151

<212> DNA

<213> Conus obscurus

<220>

<221> CDS

<222> (46)..(111)

<400> 353

tctgatggcg ggaatgccgc agcaaaagcg tttgatctaa tcaact ctg gcc ctc agg 57
 Leu Ala Leu Arg
 1

gat gaa tgc tgt gcc agt cct ccc tgt cgt ttg aat aat cca tac gta 105
 Asp Glu Cys Cys Ala Ser Pro Pro Cys Arg Leu Asn Asn Pro Tyr Val
 5 10 15 20

tgt cat tgacgacgct gatgctccag gaccctctga accacgacgt 151
 Cys His

<210> 354

<211> 22

<212> PRT

<213> Conus obscurus

<400> 354

Leu Ala Leu Arg Asp Glu Cys Cys Ala Ser Pro Pro Cys Arg Leu Asn
 1 5 10 15

Asn Pro Tyr Val Cys His
 20

<210> 355

<211> 217

<212> DNA

<213> Conus obscurus

<220>

<221> CDS

<222> (1)..(186)

<400> 355

atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
 Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

ccc act tca gat cgt gca tct gat agg agg aat gcc gca gcc aaa gcg 96
 Pro Thr Ser Asp Arg Ala Ser Asp Arg Arg Asn Ala Ala Ala Lys Ala
 20 25 30

ttt gac ctg aga tat tcg acc gcc aag aga gga tgc tgt tcc aat cct 144
 Phe Asp Leu Arg Tyr Ser Thr Ala Lys Arg Gly Cys Cys Ser Asn Pro
 35 40 45

gtc tgt tgg cag aat aat gca gaa tac tgt cgt gaa agt ggc 186
 Val Cys Trp Gln Asn Asn Ala Glu Tyr Cys Arg Glu Ser Gly
 50 55 60

taatgctcca ggaccctctg aaccacgacg t 217

<210> 356

<211> 62

<212> PRT

<213> Conus obscurus

<400> 356

Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
 1 5 10 15

Pro Thr Ser Asp Arg Ala Ser Asp Arg Arg Asn Ala Ala Ala Lys Ala
20 25 30

Phe Asp Leu Arg Tyr Ser Thr Ala Lys Arg Gly Cys Cys Ser Asn Pro
35 40 45

Val Cys Trp Gln Asn Asn Ala Glu Tyr Cys Arg Glu Ser Gly
50 55 60

<210> 357
<211> 208
<212> DNA
<213> Conus obscurus

<220>
<221> CDS
<222> (1)..(168)

<400> 357
atg ttc acc gtg ttt ctg ttg gtt gtc ttg gca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15
ttc act tca gat cgt gca tct gat ggc ggg aat gtc gca gcg tct cac 96
Phe Thr Ser Asp Arg Ala Ser Asp Gly Gly Asn Val Ala Ala Ser His
20 25 30
ctg atc gct ctg acc atc aag gga tgc tgt tct cac cct ccc tgt gcc 144
Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser His Pro Pro Cys Ala
35 40 45
cag aat aat caa gac tat tgt ggt tgacgacgct gatgctccag gaccctctga 198
Gln Asn Asn Gln Asp Tyr Cys Gly
50 55
accacgacgt 208

<210> 358
<211> 56
<212> PRT
<213> Conus obscurus

<400> 358
Met Phe Thr Val Phe Leu Leu Val Val Leu Ala Thr Thr Val Val Ser
1 5 10 15
Phe Thr Ser Asp Arg Ala Ser Asp Gly Gly Asn Val Ala Ala Ser His
20 25 30
Leu Ile Ala Leu Thr Ile Lys Gly Cys Cys Ser His Pro Pro Cys Ala
35 40 45
Gln Asn Asn Gln Asp Tyr Cys Gly
50 55

<210> 359
<211> 217
<212> DNA
<213> Conus obscurus

<220>

<221> CDS

<222> (1)..(186)

<400> 359

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atg ttc acc gtg ttt ctg ttg gtt gtc tta tca acc acc gtc gtt tcc 48
Met Phe Thr Val Phe Leu Leu Val Val Leu Ser Thr Thr Val Val Ser
  1             5             10             15

tcc act tca gat cgt gca tct gat agg agg aat gcc gca gcc aaa gcg 96
Ser Thr Ser Asp Arg Ala Ser Asp Arg Arg Asn Ala Ala Ala Lys Ala
          20             25             30

tct gac ctg atg tat tcg acc gtc aag aaa gga tgt tgt tcc cat cct 144
Ser Asp Leu Met Tyr Ser Thr Val Lys Lys Gly Cys Cys Ser His Pro
          35             40             45

gcc tgt tcg ggg aat aat cga gaa tat tgt cgt gaa agt ggc 186
Ala Cys Ser Gly Asn Asn Arg Glu Tyr Cys Arg Glu Ser Gly
          50             55             60

taatgctcca ggaccctctg aaccacgacg t 217

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<210> 360

<211> 62

<212> PRT

<213> Conus obscurus

<400> 360

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Met Phe Thr Val Phe Leu Leu Val Val Leu Ser Thr Thr Val Val Ser
  1             5             10             15

Ser Thr Ser Asp Arg Ala Ser Asp Arg Arg Asn Ala Ala Ala Lys Ala
          20             25             30

Ser Asp Leu Met Tyr Ser Thr Val Lys Lys Gly Cys Cys Ser His Pro
          35             40             45

Ala Cys Ser Gly Asn Asn Arg Glu Tyr Cys Arg Glu Ser Gly
          50             55             60

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<210> 361

<211> 157

<212> DNA

<213> Conus omaria

<220>

<221> CDS

<222> (52)..(126)

<400> 361

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ttt gat ggc ggaatgcctc agccgacagc aaagtggctg cccggatcgc t cag atc 57
                                     Gln Ile
                                     1

gac agg gat cca tgc tgt tcc tat cct gac tgt ggc gcg aat cat cca 105
Asp Arg Asp Pro Cys Cys Ser Tyr Pro Asp Cys Gly Ala Asn His Pro
          5             10             15

gag att tgt ggt gga aaa cgc tgatgctcca ggaccctctg aaccacgacg t 157
Glu Ile Cys Gly Gly Lys Arg
          20             25

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<210> 362
 <211> 25
 <212> PRT
 <213> Conus omaria

<400> 362
 Gln Ile Asp Arg Asp Pro Cys Cys Ser Tyr Pro Asp Cys Gly Ala Asn
 1 5 10 15
 His Pro Glu Ile Cys Gly Gly Lys Arg
 20 25

<210> 363
 <211> 128
 <212> DNA
 <213> Conus omaria

<220>
 <221> CDS
 <222> (26)..(88)

<400> 363
 tctcatggca ggaatgccgc acgct ctg acc gtc agg gaa tgc tgt tct cag 52
 Leu Thr Val Arg Glu Cys Cys Ser Gln
 1 5
 cct cct tgt cgc tgg aaa cat cca gaa ctt tgt agt tgaagacgct 98
 Pro Pro Cys Arg Trp Lys His Pro Glu Leu Cys Ser
 10 15 20
 gatgctccag gaccctctga accacgacgt 128

<210> 364
 <211> 21
 <212> PRT
 <213> Conus omaria

<400> 364
 Leu Thr Val Arg Glu Cys Cys Ser Gln Pro Pro Cys Arg Trp Lys His
 1 5 10 15
 Pro Glu Leu Cys Ser
 20

<210> 365
 <211> 154
 <212> DNA
 <213> Conus omaria

<220>
 <221> CDS
 <222> (52)..(123)

<400> 365
 tttgatggca ggaatgctgc agccagcgac aaagcgtctg agctgatggc t ctg gcc 57
 Leu Ala
 1
 gtc agg gga tgc tgt tcc cat cct gcc tgt gct ggg aat aat cca cat 105
 Val Arg Gly Cys Cys Ser His Pro Ala Cys Ala Gly Asn Asn Pro His
 5 10 15

atc tgt ggc aga aga cgc tgatgctcca ggaccctctg aaccacgacg t 154
 Ile Cys Gly Arg Arg Arg
 20

<210> 366
 <211> 24
 <212> PRT
 <213> Conus omaria

<400> 366
 Leu Ala Val Arg Gly Cys Cys Ser His Pro Ala Cys Ala Gly Asn Asn
 1 5 10 15

Pro His Ile Cys Gly Arg Arg Arg
 20

<210> 367
 <211> 142
 <212> DNA
 <213> Conus omaria

<220>
 <221> CDS
 <222> (40)..(102)

<400> 367
 tctgggtgtca ggaaagacgc agcgcttgcc ctgacgct ctg acc atc aag gga 54
 Leu Thr Ile Lys Gly
 1 5

tgc tgt tct gat cct agc tgt aac gtg aat aat cca gac tat tgt ggt 102
 Cys Cys Ser Asp Pro Ser Cys Asn Val Asn Asn Pro Asp Tyr Cys Gly
 10 15 20

tgacgacgct gatgctccag gaccctctga accacgacgt 142

<210> 368
 <211> 21
 <212> PRT
 <213> Conus omaria

<400> 368
 Leu Thr Ile Lys Gly Cys Cys Ser Asp Pro Ser Cys Asn Val Asn Asn
 1 5 10 15

Pro Asp Tyr Cys Gly
 20

<210> 369
 <211> 157
 <212> DNA
 <213> Conus omaria

<220>
 <221> CDS
 <222> (52)..(117)

<400> 369
 tctaattggca ggaatgccgc agccaaattc aaagcgctg cctgatgga g ctg acc 57
 Leu Thr

1

gtc agg gaa gaa tgc tgt tca gac cct cgc tgt tcc gtg gga cat caa 105
 Val Arg Glu Glu Cys Cys Ser Asp Pro Arg Cys Ser Val Gly His Gln
 5 10 15

gat atg tgt cgg tgaagcacgt gatgctccag gaccctctga accacgacgt 157
 Asp Met Cys Arg
 20

<210> 370

<211> 22

<212> PRT

<213> Conus omaria

<400> 370

Leu Thr Val Arg Glu Glu Cys Cys Ser Asp Pro Arg Cys Ser Val Gly
 1 5 10 15

His Gln Asp Met Cys Arg
 20

<210> 371

<211> 151

<212> DNA

<213> Conus purpurascens

<220>

<221> CDS

<222> (1)..(120)

<400> 371

act gat ggc agg aat gct gca gcc ata gcg ctt gac ctg atc gct ccg 48
 Thr Asp Gly Arg Asn Ala Ala Ala Ile Ala Leu Asp Leu Ile Ala Pro
 1 5 10 15

gcc gtc agg gga gga tgc tgt tcc aat cct gcc tgt tta gtg aat cat 96
 Ala Val Arg Gly Gly Cys Cys Ser Asn Pro Ala Cys Leu Val Asn His
 20 25 30

cta gaa atg tgt ggt aaa aga cgc tgatgccccca ggaccctctg aaccacgacg 150
 Leu Glu Met Cys Gly Lys Arg Arg
 35 40

t 151

<210> 372

<211> 40

<212> PRT

<213> Conus purpurascens

<400> 372

Thr Asp Gly Arg Asn Ala Ala Ala Ile Ala Leu Asp Leu Ile Ala Pro
 1 5 10 15

Ala Val Arg Gly Gly Cys Cys Ser Asn Pro Ala Cys Leu Val Asn His
 20 25 30

Leu Glu Met Cys Gly Lys Arg Arg
 35 40

<210> 373
 <211> 160
 <212> DNA
 <213> *Conus purpurascens*

<220>
 <221> CDS
 <222> (1)..(120)

<400> 373
 tct gat ggc agg gat gcc gca gcc aac gac aaa gcg tct gac ctg atc 48
 Ser Asp Gly Arg Asp Ala Ala Ala Asn Asp Lys Ala Ser Asp Leu Ile
 1 5 10 15
 gct ctg acc gcc agg aga gat cca tgc tgt ttc aat cct gcc tgt aac 96
 Ala Leu Thr Ala Arg Arg Asp Pro Cys Cys Phe Asn Pro Ala Cys Asn
 20 25 30
 gtg aat aat cca cag att tgt ggt tgaagacgct gatgctccag gaccctctga 150
 Val Asn Asn Pro Gln Ile Cys Gly
 35 40
 accacgacgt 160

<210> 374
 <211> 40
 <212> PRT
 <213> *Conus purpurascens*

<400> 374
 Ser Asp Gly Arg Asp Ala Ala Ala Asn Asp Lys Ala Ser Asp Leu Ile
 1 5 10 15
 Ala Leu Thr Ala Arg Arg Asp Pro Cys Cys Phe Asn Pro Ala Cys Asn
 20 25 30
 Val Asn Asn Pro Gln Ile Cys Gly
 35 40

<210> 375
 <211> 151
 <212> DNA
 <213> *Conus purpurascens*

<220>
 <221> CDS
 <222> (1)..(120)

<400> 375
 tct gat ggc agg gat gct gag aaa aca ggc ttt gac acg acc att gtg 48
 Ser Asp Gly Arg Asp Ala Glu Lys Thr Gly Phe Asp Thr Thr Ile Val
 1 5 10 15
 ccg gaa gac tgc tgt tcg gat cct tcc tgt tgg agg ctg cat agt tta 96
 Pro Glu Asp Cys Cys Ser Asp Pro Ser Cys Trp Arg Leu His Ser Leu
 20 25 30
 gct tgt act gga att gta aac cgc tgatgctcca ggaccctctg aaccacgacg 150
 Ala Cys Thr Gly Ile Val Asn Arg
 35 40

t

151

<210> 376
 <211> 40
 <212> PRT
 <213> *Conus purpurascens*

<400> 376
 Ser Asp Gly Arg Asp Ala Glu Lys Thr Gly Phe Asp Thr Thr Ile Val
 1 5 10 15
 Pro Glu Asp Cys Cys Ser Asp Pro Ser Cys Trp Arg Leu His Ser Leu
 20 25 30
 Ala Cys Thr Gly Ile Val Asn Arg
 35 40

<210> 377
 <211> 142
 <212> DNA
 <213> *Conus purpurascens*

<220>
 <221> CDS
 <222> (1)..(111)

<400> 377
 act gat ggc agg agt gct gca gcc ata gcg ttt gcc ctg atc gct ccg 48
 Thr Asp Gly Arg Ser Ala Ala Ala Ile Ala Phe Ala Leu Ile Ala Pro
 1 5 10 15
 acc gtc tgc tgt act aat cct gcc tgt ctc gtg aat aat ata cgc ttt 96
 Thr Val Cys Cys Thr Asn Pro Ala Cys Leu Val Asn Asn Ile Arg Phe
 20 25 30
 tgt ggt gga aga cgc tgatgccccg ggaccctctg aaccacgacg t 142
 Cys Gly Gly Arg Arg
 35

<210> 378
 <211> 37
 <212> PRT
 <213> *Conus purpurascens*

<400> 378
 Thr Asp Gly Arg Ser Ala Ala Ala Ile Ala Phe Ala Leu Ile Ala Pro
 1 5 10 15
 Thr Val Cys Cys Thr Asn Pro Ala Cys Leu Val Asn Asn Ile Arg Phe
 20 25 30
 Cys Gly Gly Arg Arg
 35

<210> 379
 <211> 157
 <212> DNA
 <213> *Conus regius*

<220>
 <221> CDS
 <222> (1)..(117)

<400> 379

tct gat gga aga aat gcc gca agc gac gcc aaa gcg ttt ccc cgg atc 48
 Ser Asp Gly Arg Asn Ala Ala Ser Asp Ala Lys Ala Phe Pro Arg Ile
 1 5 10 15

gct cca atc gtc agg gac gaa tgc tgt agc gat cct agg tgt cac ggg 96
 Ala Pro Ile Val Arg Asp Glu Cys Cys Ser Asp Pro Arg Cys His Gly
 20 25 30

aat aat cgg gac cac tgt gct tgaagacgct gctgctccag gaccctctga 147
 Asn Asn Arg Asp His Cys Ala
 35

accacgacgt 157

<210> 380

<211> 39

<212> PRT

<213> Conus regius

<400> 380

Ser Asp Gly Arg Asn Ala Ala Ser Asp Ala Lys Ala Phe Pro Arg Ile
 1 5 10 15

Ala Pro Ile Val Arg Asp Glu Cys Cys Ser Asp Pro Arg Cys His Gly
 20 25 30

Asn Asn Arg Asp His Cys Ala
 35

<210> 381

<211> 156

<212> DNA

<213> Conus regius

<220>

<221> CDS

<222> (1)..(117)

<400> 381

tct gat ggc agg aat acc gcg gcc gac gaa aaa gcg tcc gac ctg atc 48
 Ser Asp Gly Arg Asn Thr Ala Ala Asp Glu Lys Ala Ser Asp Leu Ile
 1 5 10 15

tct caa act gtc aag aga gat tgc tgt tcc cat cct ctc tgt aga tta 96
 Ser Gln Thr Val Lys Arg Asp Cys Cys Ser His Pro Leu Cys Arg Leu
 20 25 30

ttt gtt cca gga ctt tgt att tgaagacgct gctgctccag gaccctctga 147
 Phe Val Pro Gly Leu Cys Ile
 35

accacgact 156

<210> 382

<211> 39

<212> PRT

<213> Conus regius

<400> 382

Ser Asp Gly Arg Asn Thr Ala Ala Asp Glu Lys Ala Ser Asp Leu Ile

1 5 10 15
 Ser Gln Thr Val Lys Arg Asp Cys Cys Ser His Pro Leu Cys Arg Leu
 20 25 30
 Phe Val Pro Gly Leu Cys Ile
 35

<210> 383
 <211> 157
 <212> DNA
 <213> Conus regius

<220>
 <221> CDS
 <222> (1)..(117)

<400> 383
 tct gat ggc agg aat gcc gca gcc gac aac aaa gcg tct gac cta atc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asp Asn Lys Ala Ser Asp Leu Ile
 1 5 10 15
 gct caa atc gtc agg aga gga tgc tgt tcc cat cct gtc tgt aaa gtg 96
 Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Val Cys Lys Val
 20 25 30
 agg tat cca gac ctg tgt cgt tgaagacgct gctgctccag gacctctga 147
 Arg Tyr Pro Asp Leu Cys Arg
 35
 accacgacgt 157

<210> 384
 <211> 39
 <212> PRT
 <213> Conus regius

<400> 384
 Ser Asp Gly Arg Asn Ala Ala Ala Asp Asn Lys Ala Ser Asp Leu Ile
 1 5 10 15
 Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Val Cys Lys Val
 20 25 30
 Arg Tyr Pro Asp Leu Cys Arg
 35

<210> 385
 <211> 157
 <212> DNA
 <213> Conus regius

<220>
 <221> CDS
 <222> (1)..(117)

<400> 385
 tct gat ggc agg aat gcc gca gcc gac aac aga gcg tct gac cta atc 48
 Ser Asp Gly Arg Asn Ala Ala Ala Asp Asn Arg Ala Ser Asp Leu Ile
 1 5 10 15

gct caa atc gtc agg aga gga tgc tgt tcc cat cct gcc tgt aat gtg 96
Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Ala Cys Asn Val
20 25 30

aat aat cca cac att tgt ggt tgaagacgct gctgctccag gaccctctga 147
Asn Asn Pro His Ile Cys Gly
35

accacgacgt 157

<210> 386

<211> 39

<212> PRT

<213> Conus regius

<400> 386

Ser Asp Gly Arg Asn Ala Ala Ala Asp Asn Arg Ala Ser Asp Leu Ile
1 5 10 15

Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Ala Cys Asn Val
20 25 30

Asn Asn Pro His Ile Cys Gly
35

<210> 387

 $\langle 211 \rangle \cdot 157$

<212> DNA

<213> Conus regius

 $\langle 220 \rangle$

<221> CDS

$\langle 222 \rangle$ (1) . . (117)

<400> 387

tct	gat	ggc	agg	aat	gcc	gca	gcc	gac	aac	aaa	ccg	tct	gac	cta	atc	48
Ser	Asp	Gly	Arg	Asn	Ala	Ala	Ala	Asp	Asn	Lys	Pro	Ser	Asp	Leu	Ile	
1				5					10					15		

gct caa atc gtc agg aga gga tgc tgt tog cat cct gtc tgt aaa gtg 96
Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Val Cys Lys Val
20 25 30

agg tat tca gac atg tgt ggt tgaagacgct gctgctccag gaccctctga 147
Arg Tyr Ser Asp Met Cys Gly
35

accacgacgt . 157

<210> 388

<211> 39

<212> PRT

<213> Conus regius

<400> 388

Ser Asp Gly Arg Asn Ala Ala Ala Asp Asn Lys Pro Ser Asp Leu Ile
1 5 10 15

Ala Gln Ile Val Arg Arg Gly Cys Cys Ser His Pro Val Cys Lys Val
20 25 30

Arg Tyr Ser Asp Met Cys Gly
35

<210> 389
<211> 154
<212> DNA
<213> Conus stercusmuscarum

<220>
<221> CDS
<222> (1)..(114)

<400> 389
tct gat ggc agg aat gca gag cga cga caa agc gtc tgt cct ggt cgc 48
Ser Asp Gly Arg Asn Ala Glu Arg Arg Gln Ser Val Cys Pro Gly Arg
1 5 10 15
tct ggc ccc agg gga gga tgt tgt tcc cac cct gcc tgt aag gtg cat 96
Ser Gly Pro Arg Gly Gly Cys Cys Ser His Pro Ala Cys Lys Val His
20 25 30
ttt cca cac agt tgt ggt tgacgacgct gatgctccag gaccctctga 144
Phe Pro His Ser Cys Gly
35
accacgacgt 154

<210> 390
<211> 38
<212> PRT
<213> Conus stercusmuscarum

<400> 390
Ser Asp Gly Arg Asn Ala Glu Arg Arg Gln Ser Val Cys Pro Gly Arg
1 5 10 15
Ser Gly Pro Arg Gly Gly Cys Cys Ser His Pro Ala Cys Lys Val His
20 25 30
Phe Pro His Ser Cys Gly
35

<210> 391
<211> 145
<212> DNA
<213> Conus stercusmuscarum

<220>
<221> CDS
<222> (1)..(114)

<400> 391
tct gat ggc agg aat gcc gca gcc agc gac aga gcg tct gac gcg gcc 48
Ser Asp Gly Arg Asn Ala Ala Ala Ser Asp Arg Ala Ser Asp Ala Ala
1 5 10 15
cac cag gta tgc tgt tcc aac cct gtc tgt cac gtg gat cat cca gaa 96
His Gln Val Cys Cys Ser Asn Pro Val Cys His Val Asp His Pro Glu
20 25 30
ctt tgt cgt aga aga cgc tgatgctcca ggaccctctg aaccacgacg t 145

Leu Cys Arg Arg Arg Arg
35

<210> 392
<211> 38
<212> PRT
<213> Conus stercusmuscarum

<400> 392
Ser Asp Gly Arg Asn Ala Ala Ala Ser Asp Arg Ala Ser Asp Ala Ala
1 5 10 15
His Gln Val Cys Cys Ser Asn Pro Val Cys His Val Asp His Pro Glu
20 25 30

Leu Cys Arg Arg Arg Arg
35

<210> 393
<211> 154
<212> DNA
<213> Conus striatus

<220>
<221> CDS
<222> (1)..(123)

<400> 393
tct gat ggc agg aat gcc gcg gcc aac gac aaa gcg tct gac ctg gtc 48
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Asp Leu Val
1 5 10 15
gct ccg gcc atc agg gga tgc tgt tcc cac cct gtc tgt aac ttg agt 96
Ala Pro Ala Ile Arg Gly Cys Cys Ser His Pro Val Cys Asn Leu Ser
20 25 30
aat cca caa att tgt cgt gga aga cgc tga tgc tcca ggaccctctg 143
Asn Pro Gln Ile Cys Arg Gly Arg Arg
35 40
aaccacgacg t 154

<210> 394
<211> 41
<212> PRT
<213> Conus striatus

<400> 394
Ser Asp Gly Arg Asn Ala Ala Ala Asn Asp Lys Ala Ser Asp Leu Val
1 5 10 15
Ala Pro Ala Ile Arg Gly Cys Cys Ser His Pro Val Cys Asn Leu Ser
20 25 30
Asn Pro Gln Ile Cys Arg Gly Arg Arg
35 40

<210> 395
<211> 117
<212> DNA

<213> Conus textile

<220>

<221> CDS

<222> (1)..(114)

<400> 395

ttt	cat	ggc	agg	aat	gcc	gca	gcc	aaa	gcg	tct	ggc	ctg	gtc	ggg	ctg	48
Phe	His	Gly	Arg	Asn	Ala	Ala	Ala	Lys	Ala	Ser	Gly	Leu	Val	Gly	Leu	
1				5				10				15				

acc	gac	aag	agg	caa	gaa	tgc	tgt	tct	cat	cct	gcc	tgt	aac	gta	gat	96
Thr	Asp	Lys	Arg	Gln	Glu	Cys	Cys	Ser	His	Pro	Ala	Cys	Asn	Val	Asp	
		20						25					30			

cat	cca	gaa	att	tgt	cgt	tga										117
His	Pro	Glu	Ile	Cys	Arg											
		35														

<210> 396

<211> 38

<212> PRT

<213> Conus textile

<400> 396

Phe	His	Gly	Arg	Asn	Ala	Ala	Ala	Lys	Ala	Ser	Gly	Leu	Val	Gly	Leu
1				5				10				15			

Thr	Asp	Lys	Arg	Gln	Glu	Cys	Cys	Ser	His	Pro	Ala	Cys	Asn	Val	Asp
		20						25					30		

His	Pro	Glu	Ile	Cys	Arg										
		35													

<210> 397

<211> 151

<212> DNA

<213> Conus tulipa

<220>

<221> CDS

<222> (1)..(120)

<400> 397

act	gat	ggc	agg	agt	gct	gca	gcc	ata	gcg	ttt	gcc	ctg	atc	gct	ccg	48
Thr	Asp	Gly	Arg	Ser	Ala	Ala	Ala	Ile	Ala	Phe	Ala	Leu	Ile	Ala	Pro	
1				5				10				15				

acc	gtc	tgg	gaa	gga	tgc	tgt	tct	aat	cct	gcc	tgt	ctc	gtg	aat	cat	96
Thr	Val	Trp	Glu	Gly	Cys	Cys	Ser	Asn	Pro	Ala	Cys	Leu	Val	Asn	His	
		20						25					30			

ata	cgc	ttt	tgt	ggg	gga	aga	cgc	tgatgccccca	ggaccctctg	aaccacgacg	150
Ile	Arg	Phe	Cys	Gly	Gly	Arg	Arg				
		35				40					

t																151
---	--	--	--	--	--	--	--	--	--	--	--	--	--	--	--	-----

<210> 398

<211> 40

<212> PRT

<213> Conus tulipa

<400> 398

Thr Asp Gly Arg Ser Ala Ala Ala Ile Ala Phe Ala Leu Ile Ala Pro
1 5 10 15

Thr Val Trp Glu Gly Cys Cys Ser Asn Pro Ala Cys Leu Val Asn His
20 25 30

Ile Arg Phe Cys Gly Gly Arg Arg
35 40

<210> 399

<211> 157

<212> DNA

<213> Conus virgo

<220>

<221> CDS

<222> (1)..(117)

<400> 399

tct aat ggc atg aat gcc gca gcc atc agg aaa gcg tct gcc ctg gtg 48
Ser Asn Gly Met Asn Ala Ala Ala Ile Arg Lys Ala Ser Ala Leu Val
1 5 10 15

gct cag atc gcc cat cga gac tgc tgt gac gat cct gcc tgc acc gtg 96
Ala Gln Ile Ala His Arg Asp Cys Cys Asp Asp Pro Ala Cys Thr Val
20 25 30

aat aat cca ggc ctt tgc act tgaagatgct gctgccccag gaccctctga 147
Asn Asn Pro Gly Leu Cys Thr
35

accacgacgt 157

<210> 400

<211> 39

<212> PRT

<213> Conus virgo

<400> 400

Ser Asn Gly Met Asn Ala Ala Ala Ile Arg Lys Ala Ser Ala Leu Val
1 5 10 15

Ala Gln Ile Ala His Arg Asp Cys Cys Asp Asp Pro Ala Cys Thr Val
20 25 30

Asn Asn Pro Gly Leu Cys Thr
35

<210> 401

<211> 154

<212> DNA

<213> Conus geographus

<220>

<221> CDS

<222> (1)..(114)

<400> 401

tct gat ggc ggg aat gcc gca gca aaa gag tct gac gtg atc gct ctg 48
 Ser Asp Gly Gly Asn Ala Ala Ala Lys Glu Ser Asp Val Ile Ala Leu
 1 5 10 15

acc gtc tgg aaa tgc tgt acc att cct tcc tgt tat gag aaa aaa aaa 96
 Thr Val Trp Lys Cys Cys Thr Ile Pro Ser Cys Tyr Glu Lys Lys Lys
 20 25 30

att aaa gca tgt gtc ttt tgacgacgct gatgctccag gaccctctga 144
 Ile Lys Ala Cys Val Phe
 35

accacgacgt 154

<210> 402
 <211> 38
 <212> PRT
 <213> Conus geographus

<400> 402
 Ser Asp Gly Gly Asn Ala Ala Ala Lys Glu Ser Asp Val Ile Ala Leu
 1 5 10 15

Thr Val Trp Lys Cys Cys Thr Ile Pro Ser Cys Tyr Glu Lys Lys Lys
 20 25 30

Ile Lys Ala Cys Val Phe
 35

<210> 403
 <211> 154
 <212> DNA
 <213> Conus regius

<220>
 <221> CDS
 <222> (1)..(114)

<400> 403
 tct gat ggc gca gtc gac gac aaa gcg ttg gat cga atc gct gaa atc 48
 Ser Asp Gly Ala Val Asp Asp Lys Ala Leu Asp Arg Ile Ala Glu Ile
 1 5 10 15

gtc agg aga gga tgc tgt ggc aat cct gcc tgt agc ggc tcc tcg aaa 96
 Val Arg Arg Gly Cys Cys Gly Asn Pro Ala Cys Ser Gly Ser Ser Lys
 20 25 30

gat gca ccc tct tgt ggt tgaagacgct gctgctccag gaccctctga 144
 Asp Ala Pro Ser Cys Gly
 35

accacgacgt 154

<210> 404
 <211> 38
 <212> PRT
 <213> Conus regius

<400> 404
 Ser Asp Gly Ala Val Asp Asp Lys Ala Leu Asp Arg Ile Ala Glu Ile
 1 5 10 15

B20
 C9A

140

Val Arg Arg Gly Cys Cys Gly Asn Pro Ala Cys Ser Gly Ser Ser Lys
20 25 30

Asp Ala Pro Ser Cys Gly
35

B20
Conclude